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Command line parameters:

-MODEL-frame+ p2n.model -DEV=xlh
-Q=/abss/ABSSWEB spool/US10635908/runat 02062006_104214_10235/app_query.fasta_1
-Q=/abss/ABSSWEB spool/US10635908/runat_02062006_1-1.COOPCL=0 -LOOPEXT=0
-DB=GenEmbl -QFMT=fastap -SUFFIX=ye -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
-DOCALIGN=200 -THR SCORE=pCt -THR MAX=100 -THR MIN=0 -ALICN=15 -MODE=LOCAL
-OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=2000000000 -HOST=abss05h
-USER=US10635908 @CGN 1 1 5767 @runat 02062006 104214 10235 -NCPU=6 -ICPU=3
-USER=US10635908 @CGN 1 1 5767 @runat 02062006 104214 10235 -NCPU=6 -ICPU=3
-NO MMAP -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG -DEV TIMEOUT=120
-WARN TIMEOUT=30 -THREADS=1 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
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# ALIGNMENTS

Alignment Scores: Pred. No.: Score:	ORIGIN				source	FEATURES		JOURNAL		TITLE	AUTHORS	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	Locus	AX556872	RESULT 1
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                                           SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr
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                            TCGGGCTACTTTTCTATGGACTACTGGGGTCAAGGAACCTCAGTCACCGTCTCCTCA 1078
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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                                                        Bolhuis, R.L., Woehl, T. and Boettger, V. Method of producing recombinant antibodies Patent: WO 02063010-A 10 15-AUG-2002;
                                                                                                                                    synthetic construct
synthetic construct
other sequences; artificial sequences.
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Sequence 10 from Patent
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M.musculus mRNA for Ig heavy
X91667
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Direct Submission
Submitted (19-SEP-1995) E. Rosell-Vives, North State (19-SEP-1995) B. Rosell-Vives, North B. Rosell-Vives, N
                                                                                                                                                                                                                                                                                                         and Piulats,J.
Immunological and structural
antibodies mimicking an epit
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodent
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
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/mol_type="unassigned DNA"
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complementarity determining region; heavy chain; immunoglobulin variable region.
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/cell_type="fusion of splenocytes x HL1 friendly myeloma
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/note="complementarity-determining
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Rosell-Vives,E.
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
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SSLKSEDTALYYCARHRGRDSSGYVGYAIDYMGQGTSVTVSS"
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'db_xref="taxon:10090"
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Carceller, A., Rosell, E., Gomez, A., Adan Anti-idiotypic antibodies which induce epidermal growth factor receptor Patent: US 5969107-A 5 19-OCT-1999; Location/Qualifiers
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Sequence 9 1
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Carceller,A., Rosell,E., Gomez,A., Adan,J.
Anti-idiotypic antibodies which induce an epidermal growth factor receptor Patent: US 5969107-A 9 19-OCT-1999;
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PD 27-APR-
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PC A61K39/46,
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Somatic transgene immunization
BD260216
BD260216.1 GI:33069986
JP 2002542305-A/5.
Somatic transgene immunization and related methods Patent: JP 2002542305-A 5 10-DEC-2002; MAURIZIO ZANETTI, EUROGEN HOLDING SA
                                                               synthetic construct synthetic construct
                                          other sequences; artificial sequences.
1 (bases 1 to 422)
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IZIO ZAMETTI, EUROGEN HOLDING SA
Artificial Sequence
JP 2002542305-A/4
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A61K48/00,A61K31/711,A61K35/76,A61K38/00,A61K38/22,A61K38/36,
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27-APR-1999 US 09/300959
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A61K48/00,A61K31/711,A61K35/76,A61K38/00,A61K38/22,A61K38/36,
                          transgene immunization and related methods WO 0064488-A 5 02-NOV-2000; Maurizio (US)
Location/Qualifiers
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Sequence 6 from Patent WO0064488.
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MAURIZIO ZANETTI, EUROGEN HOLDING SA
OS Artificial Sequence
PN JP 2002542305-A/6
PD 10-DEC-2002
PF 27-APR-2000 JP 2000613478
PR 27-APR-2090 US 09/300959
PI MAURIZIO ZANETTI
PC A61K39/395,
PC A61K39/395,
PC A61R39/395,
PC A61R39/12,A61P37/04,C12N5/10,C12N15/09,AURIZIO ZANETTI
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A61K39/395,
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Search completed: June 3, 2006, 06:56:33 Job time: 3509.39 secs

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-UNITS_bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
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-OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=20000000000 -HOST=abbs05h
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-NARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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Ygapext
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ABQ81172
ABQ79691
ABK43187
                                                                                   급
                                                                                                                                                  SUMMARIES
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Abq81172 Anti-MN a
Abq79691 G250 heav
Abk43187 DNA encod
                                                                                   Description
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AAC90629 AAT91614 AAT71325 AAT91614 AAT71325 AAV1196 ABZ21199 ADL27490 ABL27490 ABN84611 ABZ8398 ADQ15505 AAQ15505 AAQ12682 ADQ248335 AAV248335 AAV24833 ABB12541 AAV04639 AAC87642 ADC43544 AAS11885 AAS11887 AAS	ABK43189 ABK43190 ADC24965 ADC51720 ADK51720 ABQ79699 AAT70808 AAT70810
Aac90629 Plasmid g Aac90629 Plasmid g Aat91314 CDNA enco Aat71325 Thyroid h Aav11906 Mouse PAF Ab221159 Variable Ad0127490 Nucleotid Abn84611 Anti-Stre Abn84611 Anti-Stre Ab258398 DNA encod Ad015507 Dhvar1-li Ab258397 DNA encod Ad015508 Murine 1B Ad24833 Pharmaceu Ad24833 Pharmaceu Ad24833 Pharmaceu Ac24833 Pharmaceu Ac24835 Pharmaceu Ac24835 Pharmaceu Ac24835 Pharmaceu Ac21642 Single ch Ad043554 Nucleotid Aa611885 CDNA enco Aa811885 CDNA enco Aa811885 CDNA enco Aa811885 CDNA enco Aa160477 Mouse ant Ac6379785 TRA-8 ant Ac6379785 TRA-8 ant Ac627436 TMEFF2#10 Ac627436 TMEFF2#10 Ac627436 TMEFF2#10 Ac64584 DNA seque Adv07642 anti-CD22 Adc04584 DNA seque Adt02530 Anti-CD22 Abx15270 Mouse mon Adt02524 Anti-CD22 Abx15270 Mouse mon Adt02524 Anti-CD22 Abx15270 Mouse mon Adt02524 Anti-CD22 Abx15270 Mouse mon	89 90 965 720

## ALIGNMENTS

ABQ81172 standard; cDNA; 357 BP.

ABQ81172;

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RESULT 1
ABO81172
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ID ABO8
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07-FEB-2002; 2002WO-EP001282
                                                                                                            15-AUG-2002.
                                                                                                                                                                                                                   WO200262972-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monoclonal antibody; G250; MN antigen; renal cell carcinoma; metastasis;
heavy chain; hybridoma; cancer; therapy; vaccine; mouse; gene; ss.
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1. .357
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note= "the CDS does not include a start or stop codon"
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Best Local Similarity:
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05-OCT-2001;
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SerGlyTyrPheSerMetAspTyrTrpGlyGlnGlyThrSerValThrValSerSer 119
                                                                                  LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg 100
                                                                                                                                          CTAGACACTGTGAAGGGCCGATTCACCATTTCAAGAGACAATGCCAAGAACACCCTGTAC
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                                                CTGCAAATGAGCAGTCTGAAGTCTGAGGACACAGCCTTGTTTTACTGTGCAAGACACCGC
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                                                                                                                                                                                                                                                                                                                                                 05-DEC-2002
                                                                                                                                                                                                                                                                                                                                                            ABQ79691;
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05-OCT-2001; 2001US-0327008P
                                                                                                                                                                                                                                07-FEB-2002; 2002WO-EP001283
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                                                                                                                                                                                       Bolhuis RL, Woehl T,
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                                                                                                                                                                                                                                                                                                       Homo sapiens.
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DB; ABB82004.
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/product= "VH G250"
/partial
                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                         site; diagnostic;
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                                                                                                                                                                                                                                                                                                                          therapeutic; radioimaging;
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Novel nucleic acid sequence encoding antigen-binding site of the heavy light chain of an antibody useful for expressing the antibody suitable the field of tumor diagnostics and therapeutics. in or

Claim 4; Fig 1; 18pp; English.

The invention relates to novel nucleic acid (I) encoding the antigenCC binding site of the heavy or light chain of an antibody comprising a
CC nucleotide sequence encoding the CDR3 region. (I) is useful for the
CC recombinant production of a polypeptide having an antigen-binding site,
CC by introducing (I) into a mammalian cell, culturing the cell in a medium
CC where an expression of the nucleic acid takes place, and obtaining the
CC expressed product from the medium and/or the cell. Before introducing (I)
CC into the cell, the nucleic acid is modified so that the modification does
CC not alter the amino acid sequence of the antigen-binding site of the
CC polypeptide to be expressed. The expressed product or
CC and antibody fragments are useful for preparing a diagnostic or
CC therapeutic agent and are coupled to a diagnostic marker or cytotoxic
CC agent. (I) is useful for expressing antibodies, e.g. chimerized
CC cantibodies, humanized antibody fragments. The expressed product can be
CC coupled to a diagnostic marker for use in in vitro diagnostic methods,
CC radioimaging procedures, or to a cytotoxic agent, e.g. a raddonuclide, or
CC a toxin such as cholera toxin or rich. The present sequence represents G250 heavy chain variable region encoding DNA

Sequence 357 BP; 86 A; 87 C; 94 G 90 T; 0 U; 0 Other;

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Percent Similarity:
Best Local Similarity:
                                  Alignment Scores:
Pred. No.:
1.12e-58
617.00
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Matches:
Conservative:
Mismatches:
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RESULT 3
ABK43187
ID ABK4
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The invention describes a modified virus comprising non-native polypeptides which has framework moieties each containing binding moieties, where the virus has altered tropism conferred by the binding moieties. The polypeptides can be expressed in the cytoplasm and nucleus of mammalian host cell in conformation which is maintained in absence of
                                                                                                                                                                                                                                                                                                                                                                                                                    Modified virus; adenovirus; cytostatic; gene therapy; tumour proliferating cell; cancer; vascular disease; inflammatory dinfectious disease; human immunodeficiency virus; HIV; gene;
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                                                                                                                                                                                                                                                                                                                                                                                       Human adenovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA encoding
                                                                                                      Novel modified virus comprising non-native polypeptides with stable conformation and having framework moieties containing binding moieties which confer upon the virus, an altered tropism, useful in gene therapy.
                                                                                                                                                                                                                              (GOTA-)
(GARD/)
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                                                                               Example 3;
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DB; AAU87031.
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                                                                              Page 95-99; 163pp; English.
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Percent Similarity:
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Query Match:
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                                                                                                                                                                                                          ligands for the binding moieties, where the conformation allows the binding moiety subsequently to bind with the ligand. The modified virus is useful in therapy for the preparation of a medicament for treating tumour cells, cancer, proliferating cells, vascular diseases, inflammatory diseases and infectious diseases such as Human immunodeficiency virus (HIV). The altered tropisms allow the virus to be used in treatment of disease in human or animal subjects, either by in vivo treatment of, or ex vivo treatment of cells of, the subject requiring treatment. The problems associated with the expression of functional non-native viral components in the nucleus and cytosol of host cells is solved by using the modified virus for the purpose. This sequence encodes a protein sequence used in the creation of the modified
                                                                                                                                                      Sequence 1056 BP;
                                                                                                                                                                                                virus containing non-native polypeptides
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            Conservative: Mismatches: Indels:
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Matches:
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119
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## US-10-635-908-6 101 556 496 436 376 919 81 61 41 21 LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg LeuAspThrValLysGlyArgPheThrIleSerArgAspAsnAlaLysAsnThrLeuTyr CCAGAGAAGAGGCTGGAGTTGGTCGCAGCCATTAATAGTGATGGTGGTATCACCTACTAT ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr TCCTGTGCAGCCTCTGGATTCACTTTCAGTAACTATTACATGTCTTGGGTTCGCCAGACT SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr CTGCAAATGAGCAGTCTGAAGTCTGAGGACACAGCCTTGTTTTACTGTGCAAGACACCGC CTAGACACTGTGAAGGGCCGATTCACCATTTCAAGAGACAATGCCAAGAACACCCTGTAC (1-119) x ABK43187 (1-1056)119 732 100 40 435 80 60 495 20 675 615 555

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Gaps:

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RESULT 4
ABK43189
ID ABK4
XX
ABK4
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DT 05-C
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DNA
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Modi
KW proil
KW proi
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DNA encoding fiber construct A1 G250.

05-JUN-2002

(first

entry)

proliferating cell; infectious disease; Modified virus; adenovirus; cytostatic; gene therapy; tumour cell; proliferating cell; cancer; vascular disease; inflammatory disease infectious disease; human immunodeficiency virus; HIV; gene; ds. disease; ne; ds.

Human adenovirus type

WO200208263-A2

31-JAN-2002

19-JUL-2001; 2001WO-GB003252

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RESULT 5
ABK43190
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC polypeptides which has framework moieties each containing binding CC moieties, where the virus has altered tropism conferred by the binding CC moieties, where the virus has altered tropism conferred by the binding CC moieties. The polypeptides can be expressed in the cytoplasm and nucleus of mammalian host cell in conformation which is maintained in absence of CC ligands for the binding moieties, where the conformation allows the CC binding moiety subsequently to bind with the ligand. The modified virus is useful in therapy for the preparation of a medicament for treating CC tumour cells, cancer, proliferating cells, vascular diseases transur cells, cancer, proliferating cells, vascular diseases with the virus to be cused in treatment of disease in human or animal subjects, either by in CC used in treatment of, or ex vivo treatment of cells of, the subject requiring treatment. The problems associated with the expression of C functional non-native viral components in the nucleus and cytosol of host cells is solved by using the modified virus for the purpose. This sequence encodes a protein sequence used in the creation of the modified C virus containing non-native polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel modified virus comprising non-native polypeptides with stable conformation and having framework moieties containing binding moieties which confer upon the virus, an altered tropism, useful in gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-217049/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention describes a modified virus comprising non-native
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GOTA-)
(GARD/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUL-2000; 2000GB-00017720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1411 BP; 363 A; 391 C; 349 G; 308 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; Page 104-109; 163pp; English.
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                                                          1022
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 722 GACGTGAAGCTCGTGGAGTCTGGGGGAGGCTTAGTGAAGCTTGGAGGGTCCCTGAAACTC
                                                                                                                                                                       81
                                                                                                                                                                                                                                                    61
                                                                                                                                                                                                                                                                                                                           41 ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GOT-A-GENE AB
                                                                                                                                                                                                                                                                                                                                                                                              SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr
                                                                                                                                                                 LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg 100
                                                                                                                                                                                                                                                  LeuAspThrValLysGlyArgPheThrIleSerArgAspAsnAlaLysAsnThrLeuTyr
                                                          TCGGGCTACTTTTCTATGGACTACTGGGGTCAAGGAACCTCAGTCACCGTCTCCTCA 1078
                                                                                           SerGlyTyrPheSerMetAspTyrTrpGlyGlnGlyThrSerValThrValSerSer 119
                                                                                                                                                                                                              CTAGACACTGTGAAGGGCCGATTCACCATTTCAAGAGACAATGCCAAGAACACCCTGTAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AspValLysLeuValGluSerGlyGlyGlyLeuValLysLeuGlyGlySerLeuLysLeu
                                                                                                                                   CTGCAAATGAGCAGTCTGAAGTCTGAGGACACAGCCTTGTTTTACTGTGCAAGACACCGC
                                                                                                                                                                                                                                                                                             CCAGAGAAGAGGCTGGAGTTGGTCGCAGCCATTAATAGTGATGGTGGTATCACCTACTAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5.65e-58
617.00
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100.0%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (1-1411)
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Matches:
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Indels:
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중 음 성

1013 GACGTGAAGCTCGTGGAGTCTGGGGGAGGCTTAGTGAAGCTTGGAGGGTCCCTGAAACTC

1072

AspValLysLeuValGluSerGlyGlyGlyLeuValLysLeuGlyGlySerLeuLysLeu

21 SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr 40

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The invention describes a modified virus comprising non-native conjugations which has framework moieties each containing binding consisties, where the virus has altered tropism conferred by the binding concerned by the binding consisties. The polypeptides can be expressed in the cytoplasm and nucleus of mammalian host cell in conformation which is maintained in absence of ligands for the binding moleties, where the conformation allows the confiding moiety subsequently to bind with the ligand. The modified virus is useful in therapy for the preparation of a medicament for treating cummondeficiency virus (HIV). The altered tropisms allow the virus to be used in treatment of disease in human or animal subjects, either by in convolved in treatment. The problems associated with the expression of cultivoral mon-native viral components in the nucleus and cytosol of host cells is solved by using the modified virus for the purpose. This equence encodes a protein sequence used in the creation of the modified of virus connaining non-native policiens.
US-10-635-908-6 (1-119) x ABK43190
                                                                                              Score:
                                               Query Match:
                                                              Best Local Similarity:
                                                                              Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel modified virus comprising non-native polypeptides with stable conformation and having framework moieties containing binding moieties which confer upon the virus, an altered tropism, useful in gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Modified virus; adenovirus; cytostatic; gene therapy; tumour cell; proliferating cell; cancer; vascular disease; inflammatory disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-JUN-2002
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                                                                                                                                                            Sequence 1702 BP; 447 A; 494 C; 404 G; 357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JUL-2000; 2000GB-00017720.
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                                                                                                              No.:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P-PSDB; AAU87034.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-217049/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-JUL-2001; 2001WO-GB003252
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           infectious disease; human immunodeficiency virus; HIV; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GARD/)
                                                                                                                                                                                            containing non-native polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GOT-A-GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GARDNER
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100.0%
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 (1-1702)
                                                              Length:
Matches:
Conservative:
Mismatches:
                              Gaps:
                                               Indels:
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119
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ARESULT 6
ARESULT 6
ARC 24965
AC ADC;
XX AC ADC;
XX 18-1
DT 18-1
XX MW fus;
XX MW ant;
XX W ant;
XX W ant;
XX W Syn;
XX W Sen;
XX W 21-,
XX Mus
XX Mu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          fusion protein; chimerised antibody; tumour necrosis factor; TNF; antibody; chimeric fusion protein; antiinflammatory; antiinflammatory; antiinflammatory; antiinflammatory; antiinflammatory; antiperotic; inflammatory condition; psoriasis; asthma; inflammatory bowel disease; multiple sclerosis; viral pneumonia; bacterial pneumonia; septic shock; nephritis; arteriosclerosis; mouse;
The present invention describes an isolated nucleic acid molecule, which encodes a fusion protein comprising a chimerised antibody molecule, and a tumour necrosis factor (TNF) molecule or its fragment. The antibody specifically binds to a traget. The fragment of the TNF molecule possesses the cell killing properties of a full-length TNF molecule. Also described: (1) a chimeric fusion protein (I) encoded by the isolated nucleic acid molecule; (2) an expression vector comprising the isolated nucleic acid molecule operably linked to a promoter; and (3) a recombinant cell comprising the isolated nucleic acid molecule acid molecule or the expression vector. (I) has antiinflammatory, antirheumatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New nucleic acid, useful for producing humanized or chimeric antibodies, or fusion proteins comprising the antibodies, which are G250-specific an useful for treating inflammatory conditions, e.g. rheumatoid arthritis o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-AUG-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-FEB-2002; 2002US-0355838P.
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                                                                                                                                                                                                                                                                                                                                                                                  Example 16; SEQ ID NO 50; 80pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Scott A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            heavy chain variable region DNA SEQ
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                                                                                                                    CTGCAAATGAGCAGTCTGAAGTCTGAGGACACAGCCTTGTTTTACTGTGCAAGACACCGC
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Murine G250 variable heavy chain region CDNA.

entry)

antiinflammatory; anti
inflammatory disease; granulocyte macrophage colony stimulating factor; GM-CSF; antibody; antiinflammatory; antiarthritic; antirheumatic; rhaumatoid arthritis; mouse; G250; gene; ss.

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WO2003068920-A2

21-AUG-2003

12-FEB-2003; 2003WO-US004185

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Alignment
Pred. No.:
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RESULT 8
ABQ79699
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to an isolated nucleic acid encoding a chimerised granulocyte macrophage colony stimulating factor (GM-CSF) specific antibody light chain or heavy chain. The invention further relates to: a chimerised granulocyte macrophage colony stimulating factor (GM-CSF) specific antibody light chain encoded by nucleotides 1357-1752 of a defined sequence of 6159 bp concatenated to the amino acid sequence encoded by nucleotides 1886-2203 of the 6159bp sequence; or a chimerised GM-CSF specific antibody heavy chain encoded by nucleotides 1357-1764 of a defined 6629 bp sequence concatenated to the amino acid sequence encoded by nucleotides 1839-2825 of the 6629bp sequence. The GM-CSF antibodies have antinflammatory, antiarthritic, and antirheumatic actid is useful for blocking GM-CSF and its effects on cells, such as in preventing the development of rheumatoid arthritis and other inflammatory diseases and conditions. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acids encoding chimeric granulocyte macrophage colony stimulating factor (GM-CSF) specific antibodies comprising a light and a heavy chain, useful for blocking effects of GM-CSF on cells, e.g. in preventing rheumatoid arthritis.
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ABQ79699 standard; DNA; 2431 BP
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                                                                                        SerGlyTyrPheSerMetAspTyrTrpGlyGlnGlyThrSerValThrValSerSer 119
                                                                                                                                                      LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg 100
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                                                                   TCAGGCTACTTTTCTATGGACTACTGGGGTCAAGGAACCTCAGTCACCGTCTCCTCA 431
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein of the invention.
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AspvallysLeuValGluSerGlyGlyGlyLeuValLysLeuGlyGlySerLeuLysLeu

TCCTGTGCAGCCTCTGGATTCACTTTCAGTAACTATTACATGTCTTGGGGTTCGCCAGACT SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr

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Percent Similarity:
Best Local Similarity:
Query Match:
                                                                                                                                                                                                           The invention relates to novel nucleic acid (I) encoding the antigen-
CC binding site of the heavy or light chain of an antibody comprising a
CC nucleotide sequence encoding the CDR3 region. (I) is useful for the
CC recombinant production of a polypeptide having an antigen-binding site,
CC by introducing (I) into a mammalian cell, culturing the cell in a medium
CC where an expression of the nucleic acid takes place, and obtaining the
CC expressed product from the medium and/or the cell. Before introducing (I)
CC into the cell, the nucleic acid is modified so that the modification does
CC not alter the amino acid sequence of the antigen-binding site of the
CC polypeptide to be expressed. The expressed product preferably antibodies
CC and antibody fragments are useful for preparing a diagnostic or therapeutic agent and are coupled to a diagnostic marker or cytotoxic
CC agent. (I) is useful for expressing antibodies, e.g. chimerized
CC catin antibodies, humanized antibodies, heterobispecific antibodies, and single
CC coupled to a diagnostic marker for use in in vitro diagnostic methods,
CC radioimaging procedures, or to a cytotoxic agent, e.g. a radionuclide, or
CC atoxin such as cholera toxin or ricin. The present sequence represents
CC corression vector)
US-10-635-908-6 (1-119) x ABQ79699
                                                                                                    Score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel nucleic acid sequence encoding antigen-binding site of the heavy light chain of an antibody useful for expressing the antibody suitable the field of tumor diagnostics and therapeutics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-FEB-2001; 2001US-0266853P.
05-OCT-2001; 2001US-0327008P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antibody; antigen-binding site; diagnostic; therapeutic; radioimaging; cytotoxic; G250; ds.
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RESULT 9
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                                            Murine monoclonal anti-idiotypic antibodies, 15H8, 3B6 and 5A6 are new. They induce an immune response against epidermal growth factor receptor (EGFR). The sequences of the heavy and light chain variable regions of these antibodies are given in the specification. The antibodies are used for the manufacture of drugs directed against tumours that express EGFR on their surface, including melanomas, gliomas and carcinomas
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                                                                                                                                                                                                                                                                          Monoclonal anti-idiotypic antibodies mimicking receptor - useful for tumour therapy.
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P-PSDB; AAW19577.
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                                                                                                                                                                                                                             Claim 6;
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1. .30
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/note= "no stop codon given"
1. .57
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Sequence 474

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                  26-MAY-1995;
                                                      04-DEC-1996
                                                                      EP745612-A1
                                                                                               primer_bind
                                                                                                                 primer_bind
                                                                                                                                  sig_peptide
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                                                                                                                                                                                          Mus musculus.
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cancer; neoplasia; glioma;
                                                                                                                                                                                                                                   Mouse anti-idiotypic antibody 5A6 heavy chain variable
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(MERE ) MERCK PATENT GMBH
                                    14-MAY-1996;
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1..57
/*tag= b
1..25
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                                    96EP-00107651
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Alignment Scores:
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Query Match:
DB:
RESULT 11
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ID AAC90
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AC AAC90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Murine monoclonal anti-idiotypic antibodies, 15H8, 3B6 and 5A6 are new. They induce an immune response against epidermal growth factor receptor (EGFR). The sequences of the heavy and light chain variable regions of these antibodies are given in the specification. The antibodies are used for the manufacture of drugs directed against tumours that express EGFR on their surface, including melanomas, gliomas and carcinomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Monoclonal anti-idiotypic antibodies receptor - useful for tumour therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-013659/02
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                                    Somatic transgene immunisation; immunity; disease treatment; dis
             Synthetic
                                                                         Plasmid gammalWT-TAC sequence.
                                                                                                                           AAC90629;
                                                                                                                                                     AAC90629 standard;
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                                                                                                                                                                                                                                                      GGGAGGGACAGCTCGGGCTACGTAGGGTATGCTATAGACTACTGGGGTCAAGGAACCTCA
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                                    on; immune response; disease prevention;
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RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 422 BP; 103 A; 98 C; 110 G; 111 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunization by administering ex vivo a transgene having a hematopoietic cell-specific expression element operationally linked to a polynucleotide encoding one or more heterologous epitopes to a lymphoid cell.
                                                                                                 AAT91614
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                                          22-DEC-1997
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                                                                                                                                                                                                                                LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg
                                                                                                                                                                                                                                                                                                                                             ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr
                                                                                                                                                                                                                                                                                                                                                                                                                                        GACGTGAAGCTGGTGGAGTCTGGGGGGAGGCTTAGTGAAGCTTGGAGGGTCCCTGAAACTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                         AspValLysLeuValGluSerGlyGlyGlyLeuValLysLeuGlyGlySerLeuLysLeu
                                                                                                 standard;
                                                                                                                                                                       SerGlyTyrPheSerMetAspTyrTrpGlyGlnGlyThrSerValThrValSerSer 119
                                                                                                                                                                                                               CTGCAAATGAGCAGTCTGAAGTCTGAGGACACAGCCTTGTATTACTGTGCAAGAAAGGTA
                                                                                                                                                                                                                                                                                                                          CCAGAGAAGAGGCTGGAGTTGGTCGCAGCCATTAATAGTAATGGTGGTAGCACCTACTAT
                                                                                                                                                                                                                                                                                                                                                                                  TCCTGTGCAGCCTCTGGATTCACTTTCAGTAGGTATTACATGTCTTGGGTTCGCCAGACT
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                                                                                                                                                                                                                                                                                                     Best Local Similarity:
                                                                                                                                                                                                                                                                                                                  Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                         The sequences given in AAT91614-15 encode single chain Fv (scFv) antibodies which are produced in E.coli. The scFv's are derived from an anti-T3 antibody or an anti-T9130 antibody and are produced either by:

(1) expression as an inclusion body, followed by folding (i.e. by co-expression with chaperonin. Using the methods, scFv can be produced in E. coli, either in a soluble fraction or in inclusion bodies. Like the parent double chain antibodies, the scFv polypeptides are useful in immunological diagnosis or for separation and purification, but they lessen the side effects caused by antibody constant regions. The availability of scFv polypeptides contributes to fundamental research and development of diagnostic and therapeutic drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Single chain anti-gp130 ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Producing single chain Fv antibody in Escherichia coli - by expression an inclusion body, followed by protein folding or by co-expression with chaperonin as a soluble fraction.
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                                                                                                                                                                                                                                                                                                                                                                                     Sequence 735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 5-6; 9pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P-PSDB; AAW25783.
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                                                                                                                                                           SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr 40
                                                                                                                                                                                                         AspValLysLeuValGluSerGlyGlyGlyLeuValLysLeuGlyGlySerLeuLysLeu
                                                                                 CCAGAGAAGAGGCTGGAGTTGGTCGCAGCCATTAATAGTAATGGTGGTGCTACCACCTACTAT
                                                                                                  ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr
                                                                                                                                     TCCTGTGAAGCCTCTGGATTCACTTTCAGTAGTTATTACATGTCTTGGGTTCGCCAGACT
                                                                                                                                                                                           GAGGTCAAGCTGCAGGAGTCTGGGGGGAGGCTTAGTGAAGCTTGGCGGGTCCCTGAAACTC
                                             LeuAspThrValLysGlyArgPheThrIleSerArgAspAsnAlaLysAsnThrLeuTyr
 LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in Fv; scFv; antibody; E. coli; anti-T3 antibody;
antibody; inclusion body; chaperonin; ds.
                                                                                                                                                                                                                                                                                                                                                                                    BP; 165 A; 192 C;
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/transl_except= pos:385. .387,
/note= "No stop codon given"
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                                                        US-10-635-908-6 (1-119) x AAT71325
                                                                          Query Match:
DB:
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Best Local Similarity:
                                                                                                                                Alignment Scores:
                                                                                                              Score:
                                                                                                                                                                                                                                                                                                                                                                                                                                Cbs
                                                                                                                                                                   AAT71325 shows a cDNA sequence that encodes the heavy chain variable region of thyroid hormone specific T11. A hybridoma producing T11 was created and used for T11 production. Antibody fragments are useful for identifying thyroid hormones. The hybridoma can be used to prepare the antibody having various additional functions in a large amount
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Thyroid hormone; antibody; heavy chain; light chain; variable region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT71325;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT71325 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thyroid hormone specific antibody TT1 heavy chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-SEP-1997
                                                                                                                                                                                                                                                        Gene fragment encoding thyroid hormone-recognising antibody - either the heavy or light chain variable region, recognises two different thyroid
                                                                                                                                                                                                                                                                                              WPI; 1997-292465/27
                                                                                                                                                                                                                                                                                                                                   23-OCT-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hybridoma;
                                                                                                                                                   Sequence 363
                                                                                                                                                                                                                            Claim 1; Page 5-6; 7pp; Japanese
                                                                                                                                                                                                                                                 hormones
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                                                                                                                        No.:
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{\tt CysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThrPro}
                  detection; ss.
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1. .357
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/note= "no start or
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RESULT 14
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                                                                                                                                                                                                             This sequence encodes the heavy chain hypervariable region of an antibody recognising the platelet activating factor (PAF) receptor. The variable region of mouse anti-PAF receptor monoclonal antibody is useful for the preparation of chimeric antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus sp.
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                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                             Polypeptide comprising variable region of antibody recognising receptor - useful for the preparation of chimeric antibody.
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/product= "PAF receptor antibody
/protriable region"
/note= "partial coding sequence"
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  US-10-635-908-6 (1-119) x AAV11906 (1-354)
The present invention relates to sequences for murine heavy chain variable region or light chain variable region of anti-bisphenol antibody (ABZ21157-ABZ21164 and ABB98905-ABB98912). The sequences
                                                                      Gene encoding anti-bisphenol A antibody, a recombinant protein preparation, a DNA, a vector, a transformant, preparation of a recombinant protein, a kit for determining bisphenol A.
                                                                                                                       WPI; 2003-096537/09.
P-PSDB; ABB98907.
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                                               Claim 1; Page 12; 19pp; Japanese.
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                                                                                                                                                                                                                                                          JP2002253259-A.
                                                                                                                                                                                                                                                                                                                                                                                                    light chain;
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Command line parameters:

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-Q=Abbs/ABSSWEB ppool/US10635908/runat 02062006 104216 10273/app_query.fasta_1
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-DB=EST -QFMT=fastap -SUFFIX=rst -MINMATCH=0.1 -LOOPEXT=0
-UNITS=bits -START=1 -END=-1 -MATRIX:ablosum62 -TRANS-human140.cdi -LIST=45
-DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR MIN=0 -ALIGN=15 -MODE=LOCAL
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-NO_MMAP -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG -DEV_TIMEOUT=120
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## ALIGNMENTS

	COMMENT	AUTHORS TITLE JOURNAL	ORGANISM	ACCESSION VERSION KEYWORDS SOURCE	RESULT 1 BI455668 LOCUS DEFINITION
DNA Sequencing by: Incyte Genomics, Inc.  DNA Sequencing by: Incyte Genomics, Inc.  Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:  http://inage.llnl.gov  Plate: LLAM11639 row: 1 column: 16  High quality sequence stop: 780.	Contact: Robert Strausberg, Ph.D.  Email: cgapbs-r@mail.nih.gov Tissue Procurement: Lothar Hennighausen Ph.D., Robin Humphreys CDNA Library Preparation: Life Technologies, Inc. CDNA Library Preparation: Life Technologies (Inc.) CDNA Library Arrayed by The I M A G F. Consortium (ILNE)	NIH.MGC http://mgc.nci.nih.gov/. National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999)	Mus musculus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi; Muroidea; Muridae; Murinae; Mus. 1 (hases 1 to 861)	mRNA sequence. B1455668 B1455668.1 GI:15246324 EST. Mus musculus (house mouse)	BI455668 861 bp mRNA linear EST 21-AUG-2001 603173862F1 NCI_CGAP_Mam5 Mus musculus cDNA clone IMAGE:5253279 5',

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AUTHORS
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Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Lothar Hennighausen/Robin Humphreys
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NISC_ff12c05.y1 NCI_CGAP_Mam5
, mRNA sequence.
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
                                                                             NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
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NIH"
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/mol_type="mRNA"
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/clone="IMAGE:5253279"
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                    Mus musculus (house mouse)
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602096117F1 NCI_CGAP_Co24 Mus
      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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/strain="mix FVB/N, C57BL/6J"
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musculus cDNA clone IMAGE:4215844 5',
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cDNA Library Preparation: Life Technologies cDNA Library Arrayed by: The I.M.A.G.E. Consortium/LLNL DNA Sequencing by: National Institutes of Health Intramural Sequencing Center (NISC) Clone distribution: NCI-CGAP clone distribution information found through the I.M.A.G.E. Consortium/LLNL at:
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/clone lib="NCI\_CGAP\_Mam5" / Clone lib="NCI\_CGAP\_Mam5" / Clone lib="NCI\_CGAP\_Mam5" / Cloned conditionally. Primer: Oligo do Library constructed by Life Technologies. Investigators providing samples: Lothar Hennighausen/Robin Humphreys, NIH" site 1: SalI; ner: Oligo dT.

60

311

81 LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg 100 CTGCAAATGAGCAGTCTGAGGTCTGACGACACAGCCTTGTATTACTGTTCAAGACATGGT

AACTTCTACTTT-----GACTACTGGGGCCAAGGCACCACTCTCACAGTCTCCTCA

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Tissue Procurement: Jeffrey E. Green, M.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM9792 row: b column: 05
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National Institutes of Health, Mammalian Gene
Unpublished (1999)
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Sciurognathi; Muroidea; Muridae; Murinae; Mus.
1 (bases 1 to 650)
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mRNA sequence. BF581450
BF581450 GI:11655162
                                                 BF581450 774 bp mRNA linear EST 12-DEC-2000 602100853F1 NCI_CGAP_Co24 Mus musculus cDNA clone IMAGE:4224109 5',
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Location/Qualifiers
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/clone_lib="NCI_CGAP_Co24"
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/note_2: SalI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.6 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
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/mol_type="mRNA"
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Tissue Procurement: Jeffrey E. Green, M.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arzayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
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1 (bases 1 to 774)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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Mus musculus
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High quality sequence stop: 675.
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                                                              GATATATATCGTTACTATAAAGGAACGTACTACTTTGACTATTGGGGGCCAAGGCACCACT
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                                                                                 -----SerGlyTyrPheSerMetAspTyrTrpGlyGlnGlyThrSer 114
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Site_2: SalI; Cloned unidirectionally. Primer: Oligo d'Average insert size 1.6 kb. Constructed by Life Technologies. Note: this is a NCI_CGAP Library."
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/mol_type="mRNA"
/strain="FVB/N"
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EST.
Mus musculus (house mouse)
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Tissue Procurement: Jeffrey E. Green, M.D.
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National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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http://image.llnl.gov
Plate: LLAM10998 row: e column: 12
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cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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CTGCAAATGAGCAGTCTGAGGTCTGAGGACACGGCCTTGTATTACTGTGCAAGACGGGTT
                        LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg
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Location/Qualifiers
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/lab_host="DHIOB (Tl phage-resistant)"
/clone lib="NCI CGAP CO24"
/note="Organ: colon; Vector: pCMV-SPORT6; Site_1:
Site_2: Sall; Cloned unidirectionally. Primer: Ol
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Technologies. Note: this is a NCI_CGAP Library."
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National Institutes of Health, Mammalian Gene Collection (MGC)

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Jeffrey E. Green, M.D.

cDNA Library Preparation: Life Technologies, Inc.

cDNA Library Preparation: Life Technologies, Inc.

clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

http://image.llnl.gov

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High quality sequence start: 3

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Sciurognathi; Muroidea; Muridae; Murinae; Mus.
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/clone="IMAGE:4206707"
/lab_host="DHIOB (Tl phage-resistant)"
/clone lib="NCI_CGAP_CO24"
/note="Torgan: colon; Vector: pCMV-SPORT6; Site_1:
/note="Torgan: pCMV-SPO
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Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: Jeffrey E. Green, M.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Preparation: The I.N.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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602095194F1 NCI_CGAP_Co24 Mus
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
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National Institutes of Health, Mammalian Gene Collection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov
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                                                                                                              ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr
                                                                    SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr 40
                                                  rccrordcagccrcrocarricacrraciangccarorcrrococracacr 250
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                                                                                                                                                                                   (1-119) x BF579926
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                                                                                                                                                                                                                                                                                                                                                               /db_xref="taxon:10090"
/clone="IMAGE:4209551"
/clone="IMAGE:4209551"
/clone_1ib="NHIOB_(T1 phage-resistant)"
/clone_1ib="NCI_CGAP_CO24"
/note="Organ: colon; Vector: pCMV-SPORT6; Site_1: NotI;
/note_2: Sall; Cloned unidirectionally. Primer: Oligo d'Average insert size 1.6 kb. Constructed by Life
Average insert size 1.6 kb. a NCI_CGAP_Library."
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/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           strain="FVB/N"
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466.00
88.3%
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75.5%
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Indels:
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RESULT 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ms87g10.rl Soares mouse 3NbMS Mus musculus cDNA clone IMAGE:6:5' similar to gb:X14584 IG HEAVY CHAIN PRECURSOR V-III REGION (HUMAN); gb:V00821 Mouse mRNA fragment for immunoglobulin mu encoding the C-terminus (MOUSE); mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T., Geisel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B., Theising,B., Wylle,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Euarchontoglires; Glires; Roden
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Marra M/Mouse EST Project
WashU-HHMI Mouse EST Project
WashIngton University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Louis,
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314 286 1810
/clone_lib="Soares mouse 3NbMS"
/note="Vector: pT7T3D-PacI; Site_1: Not I; Site_2: Eco RI;
ist strand cDNA was primed with a Not I - oligo(dT) primer
                                                                                                                                                                                                                                         /tissue_type="Spleen"
/dev_stage="4 weeks"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                    /clone="IMAGE:618594"
                                                                                                                                                                                                                                                                                                                                                                              /strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                               'sex="male"
                                                                                                                                                                                                                                                                                                                                                           db_xref="taxon:10090"
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RESULT 9
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                                                                                                                                                                                                                                     Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eut
Mammalla; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroldea; Muridae; Murinae; Mus.
                                                                               Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
                                                                                                                                                                                                                                                                                                         BF163874
BF163874.1 GI:11044152
EST.
                                              Plate: LLAM9204 row: 1 column: High quality sequence stop: 658.
Location/Qualifiers
                                                                                                                                                                                   Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                          NIH-MGC http://mgc.nci.nih.gov/.
                                                                                                                                                                                                                                                                                                                                                       601772386F1 NCI_CGAP_Lu29
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                                                                                                                                                                                                                                                                                                                                              mRNA sequence.
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/organism="Mus musculus"
/mol_type="mRNA"
/strain="Czech II"
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76.2%
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                                                            JOURNAL COMMENT
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AUTHORS
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VERSION
KEYWORDS
SOURCE
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BF142302
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Query Match:
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601791844F1 NCI_CGAP_Lu30 Mus
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456.00
86.8%
79.3%
73.9%
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/db_xref="taxon:10000"
/clone="IMAGE:3991460"
/tissue_type="spontaneous tumor, metastatic Stem cell origin."
/lab_host="DH10B"
/clone_lib="NCI_CGAP_Lu29"
/note="Organ: lung, Vector: pcWV-SPORT6; Site_1: Sall;
Site_2: Not1; Cloned unidirectionally. Primer: Oligo (
Library constructed by Life Technologies. Investigator providing samples: Gilbert Smith, NIH"
                                                                                                                                                                                                                     to mammary.
                                                                     dŢ.
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Length:
Matches:
Conservative:
Mismatches:
Indels:
96
96
12
3
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.
dDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium
DNA Sequencing by:Incyte Genomics, Inc. Mus musculus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Bukaryota; Metazoa; Chordata; Craniata; Fures; Rodentia; Sciurognathi; Muroidea; Muridae; Murinae; Mus. NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D. bp mRNA musculus o A linear Gene Collection (MGC) EST 24-OCT-2000 IMAGE:4022459 5', (LLNL)

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REFERENCE
AUTHORS
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DA975437
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Best Local Similarity:
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Yamashita,R., Yamamoto,J., Sekine,M., Tsuritani,K., Wakaguri,H.,
Ishii,S., Sugiyama,T., Saito,K., Isono,Y., Irie,R., Kushida,N.,
                                                                                   Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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DA975437.1
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DA975437 SYNOV2 Homo
                                                                      Hominidae; Homo.
                                                                                                                                    Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr 60
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                                                                                                                                                                                                                                                                                                                                                                           CTGCAAATGACCAGTCTAAGGTCTGAGGACACGGCCATGTATTACTGTGCA-----AGG
                                                                                                                                                                                                                                                                                                                                                                                                  LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg 100
                                                   (bases 1 to 562)
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="NCI_CGAP_Lu30"
/note="Organ: lung; Vector: pCMV-SPORT6; Site 1: NotI;
/note="Organ: lung; Vector: pCMV-SPORT6; Site 1: NotI;
Site 2: SalI; transgenic model WNT-1, expression driven by
MMTV-LTR enhancer; Cloned unidirectionally. Primer: Oligo
dT. Library constructed by Life Technologies.
Investigator providing samples: Gilbert Smith, NIH"
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/lab_host="DH10B"
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/clone="IMAGE:4022459"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NEDO human cDNA project (New Energy and Industrial Technology Developmental Organization, Japan); cDNA library construction: Helix Research Institute (HRI); 5'-end one pass sequencing: HRI, Research Association for Biotechnology (RAB) and Biotechnology Center, National Institute of Technology and Evaluation; 3'-end one pass sequencing: RAB.
sequence.
DA976755
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DA976755
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Helix Research Institute
2-6-7 Kazusa-Kamatari, Kisarazu, Chiba, 292-0818,
Tel: 81-438-52-3975
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                                SYNOV2
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/note="Vector: pME18SFL3"
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/clone="SYNOV2005900"
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sapiens cDNA clone SYNOV2007540
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Email: fl]-cdna@nifty.com

NEDO human cDNA project (New Energy and Industrial Technology

NEDO human cDNA project (New Energy and Industrial Technology

Developmental Organization, Japan); cDNA library construction:

Helix Research Institute (HRI); 5'-end one pass sequencing: HRI,

Helix Research Association for Biotechnology (RAB) and Biotechnology

Contact National Institute of Technology and Evaluation; 3'-end
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1 (bases 1 to 553)

Kimura,K., Wakamatsu,A., Sekine,M., Tsuritani,K., Wakaguri,H., Yamashita,R., Yamamoto,J., Sekine,M., Tsuritani,K., Wakaguri,H., Yamashita,R., Yamamoto,J., Saito,K., Isono,Y., Irie,R., Kushida,N., Yamashita,R., Kushida,K., Yokoi,T., Kondo,H., Wagatsuma,M., Yonoyama,T., Otsuka,R., Kanda,K., Yokoi,T., Kondo,H., Wagatsuma,M., Yonoyama,T., Otsuka,R., Kanda,K., Yokoi,T., Kondo,H., Wagatsuma,M., Yokoi,T., Saito,K., Tsogai,T., and Sugano,S. Tshibashi,T., Takahashi-Fujii,A., Tanase,T., Nagai,K., Kikuchi,H., Nakai,K., Isogai,T. and Sugano,S. Diversification of Transcriptional Modulation: Large-scale Identification and Characterization of Putative Alternative
                                                                                                                                                                                                                                                                                                                                                                                            Center, National Institute of pass sequencing: RAB.
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FLJ Project (HRI Team)
Helix Research Institute
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Genome Res. 16 (1), 55-65 (2006)
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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                     GlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArgSer 101
                                                                                                                                                                                                                                                                                                          CysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThrPro
CAAATGAACAGCCTGAGAGCCGAGGACACGGCCGTGTATTACTGTGCGAGACTCACCGGT
                                                                                                                      AspThrValLysGlyArgPheThrIleSerArgAspAsnAlaLysAsnThrLeuTyrLeu
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/note="Vector: pME18
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RESULT 13
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Rimura, K., Wakamatsu, A., Suzuki, Y., Ota, T., Nishikawa, T., Yamashita, R., Yamamoto, J., Sekine, M., Tsuritani, K., Wakaguri, H., Ishii, S., Sugiyama, T., Saito, K., Isono, Y., Irie, R., Kushida, N., Yoneyama, T., Otsuka, R., Kanda, K., Yokoi, T., Kondo, H., Wagatsuma, M., Murakawa, K., Ishida, R., Ishidashi, T., Takahashi-Fujii, A., Tanase, T., Nagai, K., Kikuchi, H., Nakai, K., Isogai, T. and Sugano, S. Diversification of Transcriptional Modulation: Large-scale Identification and Characterization of Putative Alternative Promoters of Human Genes Genome Res. 16 (1), 55-65 (2006)
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                         2 ValLysLeuValGluSerGlyGlyGlyLeuValLysLeuGlyGlySerLeuLysLeuSer 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: flj-cdna@nifty.com
NEDO human cDNA project (New Energy and Industrial Technology
NEDO human cDNA project (New Energy and Industrial Technology
Developmental Organization, Japan); cDNA library construction:
Helix Research Institute (HRI); 5'-end one pass sequencing: HRI,
Research Association for Biotechnology (RAB) and Biotechnology
Center, National Institute of Technology and Evaluation; 3'-end one
pass sequencing: RAB.
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2-6-7 Kazusa-Kamatari, K
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Contact: Takao Isogai
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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                                                                                                                                                                                                                                                                                                         /clone_lib="SYNOV2"
/note="Vector: pME1
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RS Okazaki, Y., Furuno, M., Kasukawa, T., Adachi, J., Bono, H., Kondo, S., Nikaido, I., Osato, N., Saito, R., Suzuki, H., Yamanaka, I., Nikaido, I., Osato, N., Saito, R., Suzuki, H., Yamanaka, I., Nikaido, I., Osato, N., Saito, R., Suzuki, H., Yamanaka, I., Nikaido, I., Osato, N., Saito, R., Hill, D.P., Bult, C., Kiyosawa, H., Yagi, K., Tomaru, Y., Hasegawa, Y., Nogami, A., Rischonbach, C., Gojobori, T., Baldarelli, R., Hill, D.P., Bult, C., Schonbach, C., Corbani, J., Schriml, L.M., Kanapin, A., Matsuda, H., Batalov, S., Balsel, K.W., Blake, J.A., Bradt, D., Brusic, V., Batalov, S., Corothia, C., Corbani, L.E., Cousins, S., Dalla, E., Dragani, T.A., Fletcher, C.F., Forrest, A., Frazer, K.S., Gaasterland, T., Garibooldi, M., Gissi, C., Godzik, A., Gough, J., Grimmond, S., Gariboldi, M., Gissi, C., Godzik, A., Gough, J., Jarvis, E.D., Kanai, A., Kurochkin, I.V., Lee, Y., Lenhard, B., Lyons, P.A., Maglott, D.R., Kurochkin, I.V., Lee, Y., Lenhard, B., Lyons, P.A., Maglott, D.R., Kurochkin, I.V., Lee, Y., Lenhard, B., Lyons, P.A., Maglott, D.R., Numata, K., Okido, T., Pavan, W.J., Pertea, G., Pesole, G., Numata, K., Okido, T., Pavan, W.J., Pertea, G., Pesole, G., Ringwald, M., Sandelin, A., Schneider, C., Semple, C.A., Setou, M., Shimada, K., Sultana, R., Takenaka, Y., Taylor, M.S., Tessdale, R.D., Tomita, M., Vang, L., Wang, Y., Watanabe, Y., Yang, L., Yuan, Z., Zavolan, M., Zhu, Y., Zimmer, A., Carninci, P., Yang, L., Yuan, Z., Zavolan, M., Zhu, Y., Zimmer, A., Carninci, P., Yang, L., Yuan, Z., Zavolan, M., Zhu, Y., Zimmer, A., Carninci, P., Yang, L., Yuan, Z., Zavolan, M., Zhu, Y., Zimmer, A., Carninci, P., Yang, L., Yuan, Z., Zavolan, M., Zhu, Y., Zimmer, A., Carninci, P., Yang, L., Yuan, Z., Zavolan, M., Zhu, Y., Zimmer, A., Carninci, P., Yang, L., Yuan, Z., Zavolan, M., Shikawa, T., Waki, K., Kawai, D., Shibata, K., Shibata, K., Shibata, K., Shibata, K., Shibata, Y., Taylor, M., Sasaki, D., Shibata, K., Shibata, Y., Taylor, M., Sasaki, Y., Sasaki, Y., Sasaki, Y., Sasaki, Y., Sasaki, Y.,
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BY750520 RIKEN full-length enriched, activated spleen Mus musculus CDNA clone F830004G20 5', mRNA sequence.
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
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Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center(GSC), Yokohama Institute The Institute of Physical and Chemical Research (RIKEN)
                                                                                                Nature 420, 563-573 (2002)
12466851
Contact: Yoshihide Hayashizaki
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Adachi,J., Aizawa,K., Akimura,T., Arakawa,T., Carninci,P.,
Fukuda,S., Hashizume,W., Hayashida,K., Hirozane,T., Hori,F.,
Imotani,K., Ishii,Y., Itch,M., Kagawa,I., Kawai,J., Kojima,Y.,
Kondo,S., Konno,H., Koya,S., Miyazaki,A., Murata,M., Nakamura,M.,
Nomura,K., Numazaki,R., Ohno,M., Ohsato,N., Saito,R., Sakazume,N.,
Sano,H., Sasaki,D., Sato,K., Shibata,K., Shiraki,T., Tagami,M.,
Takeda,Y., Waki,K., Watahiki,A., Muramatsu,M. and Hayashizaki,Y.
Direct Submission
81 LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg 100
                                                                                                                                                                                                                                                                                                  41
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Normalization and subtraction of cap-trapper-selected cDNAs to prepare full-length cDNA libraries for rapid discovery of new genes. Genome Res. 10 (10), 1617-1630 (2000)

RIKEN integrated sequence analysis (RISA) system-384-format sequencing pipeline with 384 multicapillary sequencer. Genome Res. 10 (11), 1757-1771 (2000)
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Fax: 81-45-503-9216
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Tissues were provided by Dr. John Todd (Dept. of Medical Genetics Tissues were provided by Dr. John Todd (Dept. of Medical Genetics Wellcome Trust Centre for Molecular Mechanisms in Disease Wellcome Wellcome Deviate (MRC building Addenbrookes Hospital Cambridge) whose assistance we gratefully acknowledge.

Please visit our web site (http://genome.gsc.riken.go.jp) for
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Okazaki,Y., I
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High-efficiency full-length cDNA cloning
Meth. Enzymol. 303, 19-44 (1999)
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Osato, N.,
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, Kasukawa, T., Adachi, J., Bono, H., Kondo, S., Saito, R., Suzuki, H., Yamanaka, I., Tomaru, Y., Hasegawa, Y., Nogami, A.,
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Schonbach, C., Gojobori, T., Baldarelli, R., Hill, D.P., Bult, C., Hume, D.A., Quackenbush, J., Schriml, L.M., Kanapin, A., Matsuda, H., Batalov, S., Beisel, K.W., Blake, J.A., Bradt, D., Brusic, V., Chothia, C., Corbani, L.E., Cousins, S., Dalla, E., Dragani, T.A., Fletcher, C.F., Forrest, A., Frazer, K.S., Gassterland, T., Gariboldi, M., Gissi, C., Godzik, A., Gough, J., Grimmond, S., Winckawa, N., Jackson, I.J., Jarvis, E.D., Kanai, A., Kurochkin, I.V., Lee, Y., Lenhard, B., Lyons, P.A., Maglott, D.R., Maltais, L., Marchionni, L., McKenzie, L., Miki, H., Nagashima, T., Marchionni, L., McKenzie, L., Miki, H., Nagashima, T., Pettovsky, N., Pillai, R., Pontius, J.U., Qi, D., Ramachandran, S., Petrovsky, N., Pillai, R., Pontius, J.U., Qi, D., Ramachandran, S., Sandelin, A., Schneider, C., Semple, C.A., Setou, M., Shimada, K., Sultana, R., Takenaka, Y., Taylor, M.S., Teasdale, R.D., Tomita, M., Verardo, R., Wagner, L., Wahlestedt, C., Wang, Y., Watanabe, Y., Verardo, R., Wilming, L.G., Wynshaw-Boris, A., Yangisawa, M., Yang, I., Yuan, Z., Zavolan, M., Zhu, Y., Zimmer, A., Carninci, P., Yuan, Z., Zavolan, M., Zhu, Y., Zimmer, A., Carninci, P., Hayatsu, N., Hirozane, Kishikawa, T., Konno, H., Nakamura, M., Sakazume, N., Sato, K., Shiraki, T., Wahi, K., Kawai, J., Aizawa, K., Sakazume, N., Sato, K., Shiraki, T., Wahi, K., Kawai, J., Aizawa, K., Shinagawa, J., Wiyazaki, A., Sakai, K., Sasaki, D., Shibata, K., Kayari, S., Shiraki, Y., Sasaki, D., Shibata, K., Kayari, S., Lander, E.S., Rogers, J., Birney, E. and Hayashizaki, Y.
FANTOM Consortium
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Analysis of the mouse transcriptome based on functional annotation of 60,770 full-length cDNAs Nature 420 (6915), 563-573 (2002) 12466851

CONSRIM

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The transcriptional landscape of the mammalian genome Science 309 (5740), 1559-1563 (2005)

Katayama,S., Tomaru, Y., Kasukawa, T., Waki, K., Nakanishi, M.,

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JOURNAL
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                                                     195
                                                                                                                                                  135 GAAGTGCAGCTGGAGTCTGGGGGAGGCTTAGTGAAGCCTGGAGGGTCCCTGAAACTC 194
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Submitted (30-MAR-2004) Yoshihide Hayashizaki, The Institute of Physical and Chemical Research (RIKEN), Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center (GSC), RIKEN Yokohama Institute; 1-7-22 Suehiro-cho, Tsurumi-ku, Yokoham Kanagawa, 230-0045, Japan (E-mail:genome-res@gsc.riken.jp, URL:http://genome.gsc.riken.jp/, Tel:81-45-503-9222, Fax:81-45-503-9216)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Arakawa, T., Carninci, P., Fukuda, S., Hashizume, W., Hayashida, K., Hori, F., Iida, J., Imamura, K., Imotani, K., Itoh, M., Kanagawa, S., Kojima, M., Konno, H., Muratta, M., Nakamura, M., Ninomiya, N., Nishiyori, H., Nomura, K., Ohno, M., Sakazume, N., Sano, H., Sasaki, D., Shibata, K., Shiraki, T., Tagami, M., Tagami, Y., Waki, K., Watahiki, A., Muramatsu, M., and Hayashizaki, Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nakamura,M., Nishida,H., Yap,C.C., Suzuki,M., Kawai,J., Suzuki,H., Carninci,P., Hayashizaki,Y., Wells,C., Frith,M., Ravasi,T., Pang,K.C., Hallinan,J., Mattick,J., Hume,D.A., Lipovich,L., Batalov,S., Engstrom,P.G., Mizuno,Y., Faghihi,M.A., Sandelin,A., Chalk,A.M., Mottagui-Tabar,S., Liang,Z., Lenhard,B. and Wahlestedt,C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Rike Genomic Sciences Center and Genome Science Laboratory in RIKEN.

Division of Experimental Animal Research in Riken contributed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RIKEN Genome Exploration Research Group
Antisense transcription in the mammalian transcriptome
Science 309 (5740), 1564-1566 (2005)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      prepare mouse tissues.
Tissues were provided by Dr. John Todd (Dept. of Medical Genetics
Wellcome Trust Centre for Molecular Mechanisms in Disease Wellcome
                                                                                                                                                                                  1 AspValLysLeuValGluSerGlyGlyGlyLeuValLysLeuGlyGlySerLeuLysLeu 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              assistance we gratefully acknowledge. Please visit our web site for further details.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              URL:http://genome.gsc.riken.jp/
URL:http://fantom.gsc.riken.jp/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Trust/MRC building Addenbrookes Hospital Cambridge) whose
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ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr 60
                                                                             SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr 40
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                                                     TCCTGTGCAGCCTCTGGATTCACTTTCAGTAGCTATGCCATGTCTTGGGTTCGCCAGACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="immunoglobulin heavy chain 4 (serum IgG1)
(MGD|MGI:96446 GB|BC024405, evidence: BLASTN, 99%,
match=1117)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /tissue_type="activated spleen"
/clone_lib="RIKEN full-length enriched mouse cDNA library"
1. .1565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="mRNA"
|strain="NOD"
|db_xref="FANTOM_DB:F830004G20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        clone="F830004G20"
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83.2%
76.5%
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Matches:
Conservative:
Mismatches:
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                                                                                                                                                                                                          81 LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg 100
                                                                                                                                                           61 LeuAspThrValLysGlyArgPheThrIleSerArgAspAsnAlaLysAsnThrLeuTyr 80
GGGGATTACGACAGGTTGCTTACTGGGGCCAAGGGACTCTGGTCACTGTCTCTGCA 488
                                                                    CTGCAAATGAGCCATCTGAAGTCTGAGGACACAGCCATGTATTACTGTGCAAGAGAT---
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Search completed: June 3, 2006, 09:16:33 Job time: 3003.94 secs

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8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -Q=/Abs/ABSSWEB spool/US10635908/runat_02062006_104220_10327/app_query.fasta_1
-DB=IEsued_Patents_Na -QFMT=fastap_SUFFIX=rni -MINNATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=Dits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DCCALIGN=200 -THR SCORE=pct -THR MAX=100 -THR MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
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Listing first 45 summaries
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-MODEL=frame+ p2n.model -DEV=xlh
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seq length: 2000000000
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                                                                                                                                                                                                                                                                                                    Issued_Patents_NA:*

1: /EMC_Celerra_SIDS3/ptodata/2/ina/5_COMB.seq:*

2: /EMC_Celerra_SIDS3/ptodata/2/ina/6A_COMB.seq:*

3: /EMC_Celerra_SIDS3/ptodata/2/ina/6A_COMB.seq:*

4: /EMC_Celerra_SIDS3/ptodata/2/ina/6A_COMB.seq:*

5: /EMC_Celerra_SIDS3/ptodata/2/ina/H_COMB.seq:*

6: /EMC_Celerra_SIDS3/ptodata/2/ina/H_COMB.seq:*

7: /EMC_Celerra_SIDS3/ptodata/2/ina/PCOMB.seq:*

8: /EMC_Celerra_SIDS3/ptodata/2/ina/PCOMB.seq:*

9: /EMC_Celerra_SIDS3/ptodata/2/ina/AE_COMB.seq:*

10: /EMC_Celerra_SIDS3/ptodata/2/ina/AE_COMB.seq:*
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Ygapop 10.0,
Fgapop 6.0,
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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            Sequence 5, Appli
Sequence 9, Appli
Sequence 31, Appli
Sequence 1, Appli
Sequence 17, Appl
Sequence 15, Appl
Sequence 15, Appl
                                                                                                                                                      Description
Patent No. 54
Sequence 83,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 5, Application US/08653402B Patent No. 5969107
                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: CARCEL
APPLICANT: ROSELL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                            ZIP: 22201
                                                                                                                                                                                                                                                                                             ADDRESSEE:
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GOMEZ, Alicia
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357	357	357	747	747	513	843	843	843	843	361	361	361	361	477	717	1347	411	372	804	711	783	923	351	351	351	414	375	375	375	357	357	357	357	2199	978	888
US-08-484-508-1	US-08-483-199	US-08-475-000-1	US-09-661-992B-8	US-09-661-992B-	US-09-318-786-3	US-09-490-324-1	US-09-490-15	US-09-490-070A-	US-09-025-769B-17	-09-490-324-	US-09-490-153-6	US-09-490-070A-6	US-09-025-769B-	US-08-653-402B-	US-07-956-399	0 5455030-	US-09-269-332-	US-09-518-737-	US-09-554-7	US-10-073-301A	US-08-487-283A-	PCT-US94-07659-	US-09-016-061-	US-08-791-391A-	US-09-339-922A-	US-08-379-057-1	US-08-759-804A-5	US-08-331-397B-	US-08-331-398A-5	US-09-227-693-21	US-08-759-804A-2	US-08-331-397B-2	US-08-331-398A-	US-09-661-992B-8	US-09-661-992B-9	US-09-661-992B-
e 15,	Sequence 15, Appl	e 15, App	85,	€ 87,	e 32, App	e 177, Ap	e 0, Ag	e 177, Ap	17	N	62,	62,	62, App	1,	Sequence 3, Appli	atent No. 54550	equence 57, Appl	equence 1,	ence 1,	equence 8,	equence 19, App	equence	equence 5, Appl	equence 5, App	equence 5, A	equence 13,	equence 58, App	equence 59,	equence 59, App	e 21, App	e 21, App	e 21,	e 21, App	equence 89, App	91,	99,

## ALIGNMENTS

```
APPLICANT: ADEN, Jaume
APPLICANT: PIULATS, Jaume
ITITLE OF INVENTION: Anti-idiotypic antibodies which induce an
ITITLE OF INVENTION: immune response against epidermal growth factor receptor
INUMBER OF SEQUENCES: 13
CORRESSEDINDENCE ADDRESS:
ADDRESSEE: Millen, White, Zelano & Branigan, P.C.
STREET: 2200 Clarendon Boulevard, Suite 1400
CITY: Arlington
STATE: Virginia
COUNTRY: U.S.A.
ZIP: 22201
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/653,402B
FILING DATE: 24-MAY-1996
CLASSIFICATION NUMBER: EP 95107967.2
FILING DATE: 26-MAY-1995
```

ATTORNEY/AGENT INFORMATION:
NAME: Lebovitz, Richard M.
REGISTRATION NUMBER: 37,06

```
REFERENCE/DOCKET NUMBER: 37,067
REFERENCE/DOCKET NUMBER: MERCK 1781
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-243-6333
TELEPAX: 703-243-6410
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 474 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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US-08-653-402B-9
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Best Local Similarity:
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                                                                                               Sequence 9, Application US/08653402B Patent No. 5969107
                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
ORIGINAL SOURCE:
ORGANISM: monoclonal ant:
   APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: monoclonal anti-idiotypic anti-EGFR antibody INDIVIDUAL ISOLATE: 5A6 heavy chain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  CCAGAGAGAGGCTGGAGTTTGTCGCAGCCATTAATAGTAATGGTGGTAGCACCTACTAT
CARCELLER, Ana
ROSELL, Elisabet
GOMEZ, Alicia
ADEN, Jaume
PIULATS, Jaume
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88.8%
90.6%
                                                                                                                                                                                                                                                                            -SerGlyTyr-----PheSerMetAspTyrTrpGlyGlnGlyThrSer 114
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                                                                                                                                                                                                                                                                                                                                                                                 297
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Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/653,402B
FILING DATE: 24-MAY-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 95107967.2
FILING DATE: 26-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: Lebovitz, Richard M.
REGISTRATION NUMBER: 37,067
REFERENCE/DOCKET NUMBER: MERCK 1781
TELEPHONE: 703-243-6333
TELEPHAN: 703-243-6330
TRELEPAX: 703-243-6410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 703-243-6410 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: monoclona INDIVIDUAL ISOLATE: FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FRAGMENT TYPE: N-terminal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: AN TITLE OF INVENTION: im NUMBER OF SEQUENCES: 1 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                     No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ANTI-SENSE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Millen, White, Zelano & Branigan, STREET: 2200 Clarendon Boulevard, Suite 1400 CITY: Arlington STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.A.
ZIP: 22201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 474 base pairs
                                                                                                                                          118 TCCTGTGCAGCCTCTGGATTCACTTTCAGTAACTATTACATGTCTTGGGTTCGCCAGACT
238 CCAGACACTGTGAAGGGCCGATTCACCATCTCCAGAGACAATGCCAAGAACACCCTGTAC
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                    61 LeuAspThrValLysGlyArgPheThrIleSerArgAspAsnAlaLysAsnThrLeuTyr
                                                                                          ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr
                                                                                                                                                                 SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr
                                                                                                                                                                                                                    GACGTGAAGCTCGTGGAGTCTGGGGGGAGGCTTAGTGAAGCTTGGAGGGTCCCTGAAACTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                monoclonal anti-idiotypic anti-EGFR antibody ISOLATE: 3B6 heavy chain
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Matches:
Conservative:
Mismatches:
Indels:
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RESULT 3
US-10-226-795-31
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DB:
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Best Local Similarity:
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APPLICANT: WILSON, JULIE
TITLE OF INVENTION: MONOCLONAL ANTIBODIES AND COMPLEMENTARITY-DETERMINING
TITLE OF INVENTION: REGIONS BINDING TO EBOLA GLYCOPROTEIN
TILE REFERENCE: ARMY 166
CURRENT APPLICATION NUMBER: US/10/226,795
CURRENT FILING DATE: 2002-11-18
NUMBER OF SEQ ID NOS: 40
SOFTWARE: PALENTIN Ver. 2.1
SEQ ID NO 31
LENGTH: 484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 31, Applica Patent No. 6875433 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
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426
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                                                                                                                                                                                                 LeuAspThrValLysGlyArgPheThrIleSerArgAspAsnAlaLysAsnThrLeuTyr 80
                                                                                                                                                                                                                                                                                                                                                                      GAAGTGCAGGTGGAGTCTGGGGGAGGCTTAGTGAAGCCTGGAGGGTCCCTGAAACTC
                                                                                                                    LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHis---
                                                                                                                                                                                                                                            CCGGAGAAGAGGCTGGAGTGGGTCGCATACATTAGTCGTGGTGGTGGTTACACCTACTAT 245
                                                                                                                                                                                                                                                                          ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr 60
                                                                                                                                                                                                                                                                                                         TCCTGTGCAGCCTCTGGATTCGCTTTCAGTAGCTATGACATGTCTTGGGTTCGCCAGACT
                             ValSerSer 119
                                                                            -----ArgSerGlyTyrPheSerMetAspTyrTrpGlyGlnGlyThrSerValThr 116
                                                          TATTACGGGAGTAGTCATTACTATGCTATGGACTACTGGGGTCAAGGAACCTCAGTCACC 425
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Matches:
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Indels:
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RESULT 4
US-08-672-176A-1
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: N/A
FILING DATE: N/A
FILING DATE: N/A
ATTORNEY/AGENT INFORMATION:
NAME: Silver, Robert S.
REGISTRATION NUMBER: unknown
REFERENCE/DOCKET NUMBER: E1042/
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-751-1142
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
IENUTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1, Application US/08672176A
Patent No. 5908925
GENERAL INFORMATION:
APPLICANT: Cohen, Margo P.; Shearman, Clyde W.
TITLE OF INVENTION: Genetically Engineered Immunoglobulins
TITLE OF INVENTION: Specificity for Glycated Albumin
                                                                                                                                                                                                                                    CELL TYPE: B-cel
CELL LINE: fused
ORGANELLE: N/A
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                            HYPOTHETICAL: no ANTI-SENSE: no FRAGMENT TYPE: N ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: DOS 4.0 or better SOFTWARE: Wordperfect, Version 5.2 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/672,176A FILING DATE: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: diskett
COMPUTER: PC
                                                                                                                                                                          LIBRARY: N/A
CLONE: pA717VH
POSITION IN GENOME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
MOLECULE TYPE: cDNA to mRNA
DESCRIPTION: A717 immunoglobulin heavy chain variable
DESCRIPTION: region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 5
                                                                                              UNITS: unknown FEATURE: antiqen
    PUBLICATION
                                                                                                                                                                                                                                                                                                                  STRAIN: Balb/c
INDIVIDUAL ISCLATE: i
DEVELOPMENTAL STAGE:
HAPLOTYPE: N/A
TISSUE TYPE: lymphoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: SSOR Market Street,
CITY: Philadelphia
STATE: Documents
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NAME/KEY: CDR-1; CDR-2; CDR-3
LOCATION: aa #31-35; aa#50-66; aa#99-109
IDENTIFICATION METHOD: similarity to known sequences
OTHER INFORMATION: antigen is human glycated albumin
UBLICATION INFORMATION: N/A
                                                                                                                                    CHROMOSOME/SEGMENT: unknown MAP POSITION: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19104
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                                                                         antigen binding sites
Y: CDR-1; CDR-2;
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                                                                                                                                                                                                                                         plasmid
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Best Local Similarity:
Query Match:
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US-08-553-497A-17
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                                                                                                                                                                                                                                                                                                                                             Sequence 17, Application US/08553497A Patent No. 5844093
                                                                                                       APPLICANT: KETTLEBOROUGH, C. A.
APPLICANT: BENDIG, MARY M.
APPLICANT: BUSIG, MARY M.
APPLICANT: GUSSOW, DETLEF
APPLICANT: GUSSOW, DETLEF
APPLICANT: MITJANS, FRANSESC
APPLICANT: MITJANS, FRANSESC
APPLICANT: BLASCO, FRANCESC
APPLICANT: BLASCO, FRANCESC
APPLICANT: PIULATS, JAIME
TITLE OF INVENTION: ANTI-EGFR SINGLE-CHAIN FVS ANT
TITLE OF INVENTION: ANTI-ODIES
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
PUBLICATION DATE:
RELEVANT RESIDUES IN SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PAGES:
DATE:
DOCUMENT NUMBER:
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                                              STREET: 2200 CL
CITY: ARLINGTON
STATE: VA
               COUNTRY: US
ZIP: 22201
                                                                            ADDRESSEE: MILLEN, WHITE, ZELANO & BRANIGAN, P.C. STREET: 2200 CLARENDON BLVD. SUITE 1400
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78.8%
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Matches:
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Mismatches:
Indels:
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Percent Similarity:
Best Local Similarity:
Query Match:
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APPLICATION NUMBER: WO PCT/EP95/00978
FILING DATE: 16-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 94104160.0
FILING DATE: 17-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 94118970.6
APPLICATION NUMBER: EP 94118970.6
APPLICATION NUMBER: BP 94118970.6
APPLICATION NUMBER: 33,302
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TELEFAX: 703-243-6410
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
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FRAGMENT TYPE: N
ORIGINAL SOURCE:
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COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/553,497A
FILING DATE: 17-NOV-1995
CLASSIFICATION: 530
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 717 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IMMEDIATE SOURCE:
CLONE: 4 B 2
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LOCATION:
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TISSUE TYPE: splenocytes
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STRAIN: Balb/c
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                                       181 CCAGACAGTGTGAAGGGGCGATTCACCATCTCCAGAGACAATGCCAAGAACACCCTGTAC
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81 LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg 100
                                                                             61 LeuAspThrValLysGlyArgPheThrIleSerArgAspAsnAlaLysAsnThrLeuTyr 80
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                                                                                                                                                          ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr
                                                                                                                                                                                                                         SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr 40
                                                                                                                                                                                                                                                                                      GAGGTGAAGCTGCAGGAGTCTGGGGGGAGACTTAGTGAAGCCTGGAGGGTCCCTGAAACTC
                                                                                                                       CCAGACAAGAGGCTGGAGTCTGTCGCAACCÁTTAGTAGTGGTGGTGCTTACATCTACTAT
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77.6%
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Matches:
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                                                             Percent Similarity:
Best Local Similarity:
US-10-635-908-6 (1-119)
                                               Query Match:
                                                                                                               Pred. No.:
                                                                                                                            Alignment Scores:
                                                                                                                                                              US-08-579-378A-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15, Application US/08579378A Patent No. 6210671
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 27-DEC-1995

CLASSIFICATION: 424

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/160,074

FILING DATE: 30-NOV-1993

PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/983,946

FILING DATE: 01-DEC-1992

PRIOR APPLICATION DATA:
O1-DEC-1992
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APPLICANT: Co, Ma
                                                                                                                                                                                                                                                                                                           TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
STREET: One MarketPlaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Co, Man Sung
TITLE OF INVENTION: Humanized i
TITLE OF INVENTION: L-Selectin
                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 95
FILING DATE: 19-SEP-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: . Floppy disk
                                                                                                                                                                                                                            TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: 415-326-2400
                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                         NAME: Liebescheutz, Joe O.
REGISTRATION NUMBER: 37,505
REFERENCE/DOCKET NUMBER: 11823-002220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: EP 9: FILING DATE: 17-AUG-1995
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OPERATING SYSTEM: PC-DOS/MS-DOS
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STATE: California
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                                                                                                                                                                                                                                                                          TYPE: nucleic acid
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                                                                                                                                                                                             NAME/KEY: CDS
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x US-08-579-378A-15 (1-405)
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Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score:
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TITLE OF INVENTION: PROCESS FOR THE ACTIVATION OF T-PA OF
ING AFTER GENETIC EXPRESSION IN PROKARYOTES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 5453363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO:2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 942,370
FILING DATE: 09-SEP-1992
APPLICATION NUMBER: 498,500
FILING DATE: 23-MAR-1990
APPLICATION NUMBER: 76,207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 4
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0
FILING DATE: 02-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 23-OCT-1986
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295 CTGCAAATGAGCAGTCTGAGGTCTGAGGACACGGCCATGTATTACTGTGCAAGAGACTAT 354
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                                                                                                                            360 CCGGAAAAGAGGCTGGAGTGGGTCGCAACCATTAGTGATGGTGGTAGTTACACCTACTAT
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81 LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg
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APPLICANT: Scheiflinger, Friedrich
APPLICANT: Kerschbaumer, Randolf
APPLICANT: Falkner, Falko-Guenter
APPLICANT: Dorner, Friedrich
APPLICANT: Dorner, Friedrich
APPLICANT: Baxter Aktiengesellschaft
TITLE OF INVENTION: Factor IX/Factor IXA Acti-
TITLE OF INVENTION: Derivatives
FILE REFERENCE: 20695C-005900US
CURRENT APPLICATION NUMBER: US/09/661,992B
CURRENT FILING DATE: 2000-09-14
PRIOR APPLICATION NUMBER: AT A157600
PRIOR FILING DATE: 1999-09-14
NUMBER OF SEQ ID NOS: 112
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 83
LENGTH: 747
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DB:
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Best Local Similarity:
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Patent No. 7033
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APPLICANT: Kerschbaumer, Randolf
APPLICANT: Falkner, Falko-Guenter
APPLICANT: Falkner, Falko-Guenter
APPLICANT: Dorner, Friedrich
APPLICANT: Baxter Aktiengesellschaft
TITLE OF INVENTION: Factor IX/Factor IXa Activating Antibodies and
TITLE OF INVENTION: Derivatives
FILE REFERENCE: 20695C-005900US
CURRENT APPLICATION NUMBER: US/09/661,992B
CURRENT APPLICATION NUMBER: AT A157600
PRIOR APPLICATION UNMBER: AT A157600
PRIOR FILING DATE: 1999-09-14
NUMBER OF SEQ ID NOS: 112
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Best Local Similarity:
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SEQ ID NO 99
LENGTH: 888
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LOCATION: (228)
OTHER INFORMATION: n = 9
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APPLICANT: Scheiflinger, Friedrich
APPLICANT: Kerschbaumer, Randolf
APPLICANT: Kerschbaumer, Randolf
APPLICANT: Falkner, Falko-Guenter
APPLICANT: Dorner, Friedrich
APPLICANT: Dorner, Friedrich
APPLICANT: Baxter Aktiengesellschaft
TITLE OF INVENTION: Factor IX/Factor IXA Acti
TITLE OF INVENTION: Derivatives
FILE REFERENCE: 20695C-005900US
CURRENT APPLICATION NUMBER: US/09/661,992B
CURRENT FILING DATE: 2000-09-14
PRIOR APPLICATION NUMBER: AT A157600
PRIOR FILING DATE: 1999-09-14
INUMBER OF SEG ID NOS: 112
SOFTWARE: PatentIn Ver. 2.1
SEG ID NO 91
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OTHER INFORMATION: n = 1
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APPLICANT: Scheiflinger, Friedrich
APPLICANT: Kerschbaumer, Randolf
APPLICANT: Falkner, Falko-Guenter
APPLICANT: Darner, Friedrich
APPLICANT: Dorner, Friedrich
APPLICANT: Baxter Aktiengesellschaft
TITLE OF INVENTION: Factor IX/Factor IXA Activating Antibodies and
TITLE OF INVENTION: Derivatives
FILE REFERENCE: 20695C-005900US
CURRENT APPLICATION NUMBER: US/09/661,992B
CURRENT APPLICATION NUMBER: AND ALST ALST AND ALST AND
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SEQ ID NO 89
LENGTH: 2199
TYPE: DNA
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:198A2
OTHER INFORMATION: scFv-alkaline phosphatase fusion protein
OTHER INFORMATION: expression vector pDAP2-198AB2#100)
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TCCTCA 432
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Matches:
Conservative:
Mismatches:
Indels:
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RESULT 12 US-08-31-398A-21 ; Sequence 21, Application US/08331398A ; Patent No. 5608039

GENERAL INFORMATION:

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LOCATION: 1..357
OTHER INFORMATION:
US-08-331-398A-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: HUNTEY, TOM
REGISTRATION NUMBER: 38,498
REFERENCE/DOCKET NUMBER: 0152
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9640
TELEPAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 21:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Pastan, Ira
APPLICANT: Willingham, Mark
APPLICANT: Willingham, Mark
APPLICANT: FitzGerald, David
APPLICANT: FitzGerald, Ulrich
APPLICANT: Brinkmann, Ulrich
APPLICANT: Pai, Lee
TITLE OF INVENTION: Single Chain B3 Antibody Fusion Proteins
TITLE OF INVENTION: and Their Uses (as amended)
NUMBER OF SEQUENCES: 68
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                 Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 357 base pairs
TYPE: nucleic acid
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APPLICATION NUMBER: US 07/596,289
ETLING DATE: 12-CCT-1990
ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 30-SEP-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: single
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121
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                                                                                               SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr
                             ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr
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One Market Plaza, Steuart Street Plaza
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%R: 015280-126110US
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Matches:
Conservative:
Mismatches:
Indels:
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                                 Percent Similarity:
Best Local Similarity:
                                                                    Score:
                                                                                                       Alignment Scores:
                                                                                                                                          US-08-331-397B-21
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                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 357 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/331,397B
FILING DATE: 28-OCT-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/767,331
FILING DATE: 30-SEP-1991
PRIOR APPLICATION NUMBER: US 07/596,289
ETITING DATE: 13-OCT-1890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL
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REGISTRATION NUMBER: 38,498
REGISTRATION NUMBER: 0152
REFERENCE/DOCKET NUMBER: 0152
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1155 543-9600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Pastan, I
APPLICANT: Benhar, I
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                    No . :
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                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                             MOLECULE TYPE:
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                                                                                                                                                         LOCATION: 1..357
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 12-OCT-1990
                                                                                                                                                                                              NAME/KEY:
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Specific Antibody Fragments, Fusion Proteins, and
Thereof
                                                                                                                                                         /note= "B5 Variable Heavy chain (V-H)"
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                                                                    Length:
Matches:
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                                                     Conservative:
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Query Match: DB:

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                                                            FILING DATE: 30-SEP-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/596,289
FILLING DATE: 12-OCT-1990
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen L.
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 015280-126140US
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Pastan, Ira
APPLICANT: Willingham, Mark
APPLICANT: FitzGerald, David J.
APPLICANT: Brinkmann, Ulrich
APPLICANT: Brinkmann, Ulrich
TITLE OF INVENTION: Tumor-Specific Antibody Fragments,
TITLE OF INVENTION: Fusion Proteins, and Uses Thereof
NUMBER OF SEQUENCES: 68
                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIOM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/331,398
FILING DATE: 28-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/767,331
                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 03-DEC-19
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CITY: San Francisco
                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
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                                                 TELEPHONE:
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CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
                (415) 576-0300
OR SEQ ID NO: 2
                                                     (415)
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                                                   576-0200
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Patent No. 62
                                                                                                                                                                                                                                  GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: BENHAR, Itai
APPLICANT: BENHAR, Itai
APPLICANT: PADLAN, Eduardo A.
APPLICANT: JUNG, Sun-Hee
APPLICANT: LEE, Byungkook
APPLICANT: LEE, HUMANIZED TUMOR-SPECIFIC ANTIBODY
TITLE OF INVENTION: FRAGMENTS, FUSION PROTEINS, AND USES THEREOF
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
TOWNSONDENCE ADDRESS:
                                               SOFTWARE: PATENTIN Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/227,69: FILING DATE:
                                                                                                                                                                          STREET: Steuart StreCITY: San Francisco
STATE: California
                                                                                                                                                                                                                               ADDRESSEE:
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Application US/09227693

E: Townsend and Townsend Khourie and Crew Steuart Street Tower, One Market Plaza

Tower,

US/09/227,693

Version

#1

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US-10-635-908-6 (1-119) x US-08-759-804A-21 (1-357)
                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-759-804A-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION:
301
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                                                            241
                                                                                                                      181 CCAGACACTGTAAAGGGCCGATTCACCATCTCCAGAGACACGCCAAGAACACCCTGTAC 240
                                                                                                                                                                                    121
                                                                            81 LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg 100
                                                                                                                                              61 LeuAspThrValLysGlyArgPheThrIleSerArgAspAsnAlaLysAsnThrLeuTyr
                                                                                                                                                                                                              41 ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr
                                                                                                                                                                                                                                                13
                                                                                                                                                                                                                                                                      21 SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr 40
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                    SerGlyTyrPheSerMetAspTyrTrpGlyGlnGlyThrSerValThrValSerSer 119
                                                           CTGCAGATGAGCCGTCTGAAGTCTGAGGACACAGCCATGTATTACTGTGCAAGGGGGGCTC 300
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; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: 1..357
; OTHER INFORMATION: /standard name= "B5
; OTHER INFORMATION: variable heavy chain"
US-09-227-693-21
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Best Local Similarity:
Query Match:
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APPLICATION NUMBER: 08/331,396
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/767,331
FILING DATE: 30-SEP-1991
PRIOR APPLICATION NUMBER: US 07/596,289
PILING DATE: 12-OCT-1990
ATTORNEY AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 15280-126-1-3
REFERENCE/DOCKET NUMBER: 15280-126-1-3
REFERENCE/DOCKET NUMBER: 15280-126-1-3
REJEPHONE: (415) 543-9600
TELEPHONE: (415) 543-9600
TELEPHONE: (415) 543-9600
TELEPHONE: (415) 543-9603
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 357 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: unknown
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Search completed: June 3, 2006, 06:53:41 Job time: 121.643 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Alignment Scores:
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                                                                                                                                 81 LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg 100
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241 CTGCAGATGAGCCGTCTGAAGTCTGAGGACACAGCCATGTATTACTGTGCAAGGGGGCTC 300
                                                                                                                                                                                                                         61 LeuAspThrValLysGlyArgPheThrIleSerArgAspAsnAlaLysAsnThrLeuTyr
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                                                                 TCTGATGGTTCCTGGTTTGCTTACTGGGGGCCAAGGGACTCTGGTCACTGTCTCCTCA 357
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Matches:
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Title:
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-MODEL=frame+ p2n.model -DEV=xlh
-Q=|abss|ABSSWEB spool/US10635908/runat 02062006_104222_10380/app_query.fasta_1
-Q=|abss|ABSSWEB spool/US10635908/runat 02062006_104222_10380/app_query.fasta_1
-DB=published Applications NA Main -QFMT=fastap -SUFFIX=rnpbm -MINVATCH=0.1
-DOPCL=0 -LOOPEXT=0 -UNITS=bIts -START=1 -END=-1 -MATRIX=blosum62
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100
-THR_MIN=0 -ALIGN=15 -MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
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Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Database
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                                                                                                 Result
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -USER=US10635908 @CGN 1 1 2326 @runat 02062006 104222 10380 -NCPU=6 -ICPU=3 -NO MMAP -NEG SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG -DEV TIMEOUT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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                                                                    No.
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                                                                 Score
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seq length: 2000000000
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   617
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1: | EMC_Celerra_SIDSJ/ptodata/2/pubpna/US07_PUBCOMB.seq:*

2: | EMC_Celerra_SIDSJ/ptodata/2/pubpna/US08_PUBCOMB.seq:*

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10: | / EMC_Celerra_SIDSJ/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

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12: | / EMC_Celerra_SIDSJ/ptodata/2/pubpna/US11A_PUBCOMB.seq:*

13: | / EMC_Celerra_SIDSJ/ptodata/2/pubpna/US11A_PUBCOMB.seq:*

14: | / EMC_Celerra_SIDSJ/ptodata/2/pubpna/US11A_PUBCOMB.seq:*

15: | / EMC_Celerra_SIDSJ/ptodata/2/pubpna/US11A_PUBCOMB.seq:*

16: | / EMC_Celerra_SIDSJ/ptodata/2/pubpna/US11A_PUBCOMB.seq:*

16: | / EMC_Celerra_SIDSJ/ptodata/2/pubpna/US11A_PUBCOMB.seq:*
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Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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US-10-470-940-1
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Sequence 1, Appli
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-436-783-1	1001001001	10-436-783-	0 10-10-9E9-971-	-10-461-847-	US-10-452-440-	US-10-305-231-	3 US-09-900-590-5	US-08-791-391A-	US-08-790-5	10-354-246-3	US-10-258-728	3 US-11-093-103-8	3 US-11-093-103-	-11-093-103-9	3 US-11-093-103-8	0 US-10-789-273-	-10-232-030	US-10-704-070	US-10-70	0-388-389	-10-010-942	US-09-286-240-3	5	US-10-300-215-	US-10-383-447-5	US-10-275-180A	US-10-286-132A-2	-10-281-479A-2	US-10-169-351-	US-10-169-351-6	-10-706-391-	US-10-077-62	US-0	-10-706-391-	US-10-077-624-	-09-910-358D-	US-09-881-823-	US-10-226-795-	1 US-10-504-389A	US-10-635-908-9	-10-333-235A-	US-10-333-235A-1	US-10-333-235A-8	US-10-635-908-5	
sequence 13, Appr	94401100	45 Appl	S Ann	5. Appl	5, Appl	<b>υ</b>		5	ū	Sequence 3, Appli	-	Sequence 89, App	e 91,	equence 99,	e 83	e 3,	quence 3,	quence 3,	e u	quence 3,	quence 3,	equence 3,	Sequence 1,	quence 88, Appl	equence 5, Appl	equence 21,	equence 21,	equence 21,	equence 68,	equence 66,	e 1,	equence 1,	e 1,	e 3,	ი ა	e 5	e 11,	e 31	Sequence 50,	quence 9, Appii	equence 11,	equence	е 8,	equence 5, A	

## ALIGNMENTS

US-10-470-940-1

GENERAL INFORMATION:

Sequence 1, Application US/10470940 Publication No. US20040077081A1

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CURRENT APPLICATION NUMBER: US/10/470,940
CURRENT FILING DATE: 2003-08-19
PRIOR APPLICATION NUMBER: PCT/ EPOZ/ 01282
PRIOR FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: US 60/ 266853
PRIOR FILING DATE: 2001-02-07
PRIOR APPLICATION NUMBER: US 60/ 327008
PRIOR FILING DATE: 2001-10-05
PRIOR FILING DATE: 2001-10-05
                                                                                 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Oosterwijk, Egbert
APPLICANT: Warnaar, Sven
APPLICANT: Ullrich, Stefan
TITLE OF INVENTION: Hybridoma Cell Line G250 and its use for producing monoclonal
TITLE OF INVENTION: antibodies
FILE REFERENCE: 2923-0548
                                                                                                       NUMBER OF SEQ ID NOS: 2
SOFTWARE: PatentIn version 3.2
                           TYPE: DNA
ORGANISM: Artificial Sequence
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APPLICANT: Boettger, Volker
TITLE OF INVENTION: Method of Producing Recombinant Antibodies
FILE REFERENCE: 2923-552
CURRENT APPLICATION NUMBER: US/10/635,908
CURRENT FILING DATE: 2003-08-07
PRIOR APPLICATION NUMBER: PCT/PF02/01283
PRIOR FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 60/327,008
PRIOR APPLICATION NUMBER: 60/327,008
PRIOR PRIOR DATE: 2001-0-05
PRIOR APPLICATION NUMBER: 60/266,853
PRIOR FILING DATE: 2001-02-07
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn version 3.2
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APPLICANT: Woehl, Thorsten
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FEATURE:
REATURE:
NAME/KEY: misc_feature
LOCATION: (91)..(105)
OTHER INFORMATION: sequence
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NAME/KEY: misc feature
LOCATION: (148)..(198)
OTHER INFORMATION: sequence
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Percent Similarity:
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Query Match:
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APPLICANT: GOT-A-GENE AB

APPLICANT: Leif, Lindholm

APPLICANT: Karin, Nord

APPLICANT: Rebecca, Gardner

IITLE OF INVENTION: Modified Virus

FILE REFERENCE: 9.7.72728/001

CURRENT APPLICATION NUMBER: US/10/333,235A

CURRENT FILING DATE: 2003-12-04

NUMBER OF SEQ ID NOS: 71

SOFTWARE: Patentin version 3.0

SEQ ID NO 8
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                                 Percent Similarity:
Best Local Similarity:
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                Query Match:
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LENGTH: 357
                                                                                                                                                             OTHER INFORMATION: G250 single chain antibody construct
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ORGANISM: Artificial
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Sequence 10, Application US/1033235A

Publication No. US20040132007A1

GENERAL INFORMATION:
APPLICANT: GOT-A-GENE AB
APPLICANT: Leif, Lindholm
APPLICANT: Leif, Lindholm
APPLICANT: Rebecca, Gardner
TITLE OF INVENTION: Modified Virus
FILE REFERENCE: 9.7.77278/001

CURRENT APPLICATION UNMER: US/10/333,235A

CURRENT APPLICATION UNMER: US/10/333,235A
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ORGANISM: Artificial
FEATURE:
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APPLICANT: GOTA-GENE AB
APPLICANT: Leif, Lindholm
APPLICANT: Karin, Nord
APPLICANT: Flerre, Boulanger
APPLICANT: Pierre, Boulanger
APPLICANT: Rebecca, Gardner
TITLE OF INVENTION: Modified Virus
FILE REFERENCE: 9.7,72728/001
CURRENT APPLICATION NUMBER: US/10/333,235A
CURRENT FILING DATE: 2003-12-04
NUMBER OF SEQ ID NOS: 71
NUMBER OF SEQ ID NOS: 71
SOFTWARE: Patentin version 3.0
SEQ ID NO 11
LENGTH: 1702
TYPE: DNA
ORGANISM: Artificial
ORGANISM: Artificial
FEATURE:
OTHER INFORMATION: Fiber construct A7 G250
US-10-333-235A-11
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Best Local Similarity:
Query Match:
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US-10-333-235A-11
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RESULT 6
US-10-635-908-9
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Publication No. US20040132007A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                           ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr 60
                                                        CTGCAAATGAGCAGTCTGAAGTCTGAGGACACAGCCTTGTTTTACTGTGCAAGACACCGC
                                                                                                                             LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg 100
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                                                                                                                                                                                     LeuAspThrValLysGlyArgPheThrIleSerArgAspAsnAlaLysAsnThrLeuTyr 80
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Matches:
Conservative:
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FILE REFERENCE: 2923-552
CURRENT APPLICATION NUMBER: US/10/635,908
CURRENT FILING DATE: 2003-08-07
PRIOR APPLICATION NUMBER: PCT/EP02/01283
PRIOR FILING DATE: 2002-02-07
PRIOR APPLICATION NUMBER: 60/327,008
PRIOR APPLICATION NUMBER: 60/327,008
PRIOR FILING DATE: 2001-00-05
PRIOR PRIOR DATE: 2001-00-07
PRIOR FILING DATE: 2001-03-07
NUMBER: 60/266,853
PRIOR FILING DATE: 2001-03-07
NUMBER: 00 SEQ ID NOS: 23
SOPTWARE: PATENTIN VERSION 3.2
SEQ ID NO 9
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DB:
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US-10-504-389A-50
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                                                                                                                                                                                 Sequence 50, Application US/10504389A Publication No. US20060045876A1 GENERAL INFORMATION:
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APPLICANT: Renner, Christoph
APPLICANT: Scott, Andrew
TITLE OF INVENTION: FUSION PROTEINS OF HUMANIZED
TITLE OF INVENTION: ANTIBODIES AND USES THEREOF
FILE REFERENCE: LUD 5821
CURRENT APPLICATION NUMBER: US/10/504,389A
CURRENT FILING DATE: 2004-08-10
PRIOR APPLICATION NUMBER: PCT/US03/04243
PRIOR FILING DATE: 2002-02-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Bolhuis, Reinier LH
APPLICANT: Woehl, Thorsten
APPLICANT: Boetcger, Volker
TITLE OF INVENTION: Method of Producing Recombinant Antibodies
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Conservative:
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Sequence 31, Application US/10226795

| Sequence 31, Application US20040053865A1 |
| Publication No. MARY KATE |
| APPLICANT: WILSON, JULIE |
| TITLE OF INVENTION: MONOCLONAL ANTIBODIES AND COMPLEMENTARITY-DETERMINING |
| TITLE OF INVENTION REGIONS BINDING TO EBOLA GLYCOPROTEIN |
| PILE REFERENCE: ARMY 166 |
| CURRENT APPLICATION NUMBER: US/10/226,795 |
| CURRENT APPLICATION UNMBER: US/10/226,795 |
| CURRENT FILING DATE: 2002-11-18 |
| NUMBER OF SEQ ID NOS: 40 |
| SOFTWARE: PatentIn Ver. 2.1 |
| SEQ ID NO 31 |
| LENGTH: 484 |
| TYPE: DNA |
| CONTROL OF THE STATE OF THE 
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US-10-226-795-31
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SEQ ID NO 50
; LENGTH: 444
TYPE: DNA
ORGANIAM: Mus musculus
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                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence; FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic heavy
OTHER INFORMATION: chain of Mab EGP13F6-1-2 DNA sequence
US-10-226-795-31
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Best Local Similarity:
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                                                               Percent Similarity:
Best Local Similarity:
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OTHER INFORMATION:
                                                                                                                                                                No.:
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Matches:
Conservative:
Mismatches:
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Matches:
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Query Match: DB:

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Query Match:
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                                                           US-10-635-908-6 (1-119) x US-09-881-823-11 (1-466)
                                                                                                                                                                                     Alignment Scores:
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; NAME/KEY: CDS
; LOCATION: (11)..(442)
US-09-881-823-11
                                                                                                                                                                      Pred. No.:
                                                                                                                                                                                                                                                                                                      APPLICANT: CHEN, LI
TITLE OF INVENTION: Method for the Treatment and Prevention of Dental Caries
FILE REFERENCE: 22851-032
CURRENT APPLICATION NUMBER: US/09/881,823
CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION UNMBER: US 07/378,577
PRIOR APPLICATION UNMBER: US 07/378,577
PRIOR FILING DATE: 1999-08-20
NUMBER OF SEQ ID NOS: 32
SOFTWARE: Patentin version 3.0
SEQ ID NO 11
LENGTH: 466
TYPE: NAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence 11, Application US/09881823
Patent No. US20020068066A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Murine
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ANDERSON, MAXWELL MORRISON, SHERIE
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      127
      US-10-635-908-6 (1-119)
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Sequence 5, Application US/09910358D
Publication No. US20040052814A1
GENERAL INFORMATION:
APPLICANT: Shi, Wenyuan
APPLICANT: Anderson, Maxwell
APPLICANT: Morrison, Sherie
APPLICANT: Trinh, Kham
APPLICANT: Wims, Letitia
APPLICANT: Chen, Li
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US-09-910-358D-5
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                                        Percent Similarity:
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Query Match:
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, NAME/KEY: CDS
; LOCATION: (159)..(533)
; OTHER INFORMATION: VH
US-09-910-358D-5
                                                                                                            Alignment Scores:
Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.1
SEQ ID NO 5
LENGTH: 533
TYPE: DNA
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                                                                                                                                                                                                                                          FEATURE:
NAME/KEY: CDS
LOCATION: (111)..(158)
OTHER INFORMATION: Linker Peptide
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NAME/KEY: CDS
LOCATION: (69)..(110)
OTHER INFORMATION: Dhvar
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x US-09-910-358D-5 (1-533)
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FILE REFERENCE: 2101363-991200
CURRENT APPLICATION NUMBER: US/10/077,624
CURRENT FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 09/910,358
PRIOR PILING DATE: 2001-07-19
PRIOR APPLICATION NUMBER: US 09/378,577
PRIOR PILING DATE: 1999-08-20
NUMBER OF SEQ ID NOS: 31
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 5
TYPE: DAY
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US-10-077-624-5
; Sequence 5, Application US/10077624
; Publication No. US20030143234A1
; GENERAL INFORMATION:
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                                                          US-10-635-908-6 (1-119)
                                                                                                                                                                                                                  ; OTHER INFORMATION: Synthesized using US-10-077-624-5
                                                                                                                                                                                 Alignment Scores:
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APPLICANT: Trinh, Kham
APPLICANT: Wims, Letitia
APPLICANT: Chen, Li
APPLICANT: Qi, Fengxia
TITLE OF INVENTION: ANTI-MICROBIAL TARGETING CHIMERIC PHARMACEUTICAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: THE REGENTS OF THE UNIVERSITY OF CALIFORNIA APPLICANT: WASHINGTON DENTAL SERVICE APPLICANT: Shi, Wenyuan APPLICANT: Anderson, Maxwell
                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Artificial sequence
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-----HisArgSerGlyTyrPheSerMetAspTyrTrpGlyGlnGlyThrSerValThr 116
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APPLICANT: Anderson, Maxwell H.
TITLE OF INVENTION: ANTI-MICROBIAL TARGETING CHI
FILE REFERENCES: 2101363-991600
CURRENT APPLICATION NUMBER: US/10/706,391
CURRENT FILING DATE: 2003-06-16
PRIOR APPLICATION NUMBER: US 10/077,624
PRIOR FILING DATE: 2002-02-14
PRIOR FILING DATE: 2002-02-14
PRIOR FILING DATE: 2002-07-19
PRIOR FILING DATE: 2001-07-19
PRIOR FILING DATE: 2001-07-19
PRIOR FILING DATE: 1999-08-20
NUMBER OF SEQ ID NOS: 71
SOFTWARE: Patentin version 3.1
SEQ ID NO 3
LENGTH: 533
TYPE: DNA
ROGANISM: Artificial sequence
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US-10-706-391-3
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                                                                                                                      US-10-635-908-6 (1-119) x US-10-706-391-3 (1-533)
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                                                                                                                                                                                                                                 Alignment Scores: Pred. No.:
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159 GACGTGAAGCTTGTGGAGTCTGGGGGGAGGCTTAGTGAAACCCTGGAAGGGTCCCTGAAACTC
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                                                                               AspValLysLeuValGluSerGlyGlyGlyLeuValLysLeuGlyGlySerLeuLysLeu
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Conservative:
Mismatches:
Indels:
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US-09-910-358D-1
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                                                                                          Best Local Similarity:
Query Match:
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; LOCATION: (189)..(563)
; OTHER INFORMATION: VH
US-09-910-358D-1
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                                                     US-10-635-908-6 (1-119) x US-09-910-358D-1 (1-563)
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APPLICANT: Ande
APPLICANT: Morr
APPLICANT: Trin
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CURRENT APPLICATION NUMBER: US/09/910,358D
CURRENT FILING DATE: 2001-07-19
PRIOR APPLICATION NUMBER: US 09/378,577
PRIOR FILING DATE: 199-08-20
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Fusion Proteins for Targeted Delivery of Antimicrobial Peptides FILE REFERENCE: 22851-033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Wims, Letitia APPLICANT: Chen, Li
                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 563
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                        NAME/KEY: CDS
LOCATION: (141)..(188)
OTHER INFORMATION: Linker Peptide
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LOCATION: (69)..(140)
OTHER INFORMATION: Hi
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Morrison, Sherie
GACGTGAAGCTTGTGGAGTCTGGGGGGAGGCTTAGTGAACCCTGGAGGGTCCCTGAAACTC 248
               AspValLysLeuValGluSerGlyGlyGlyLeuValLysLeuGlyGlySerLeuLysLeu
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                                                                                                                                                                                                                                                                                                                      SEQ ID NO 1
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                                         189 GACGTGAAGCTTGTGGAGTCTGGGGGGAGGCTTAGTGAACCCCTGGAGGGTCCCTGAAACTC
21 SerCysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThr 40
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Percent Similarity:
Best Local Similarity:
Query Match:
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TITLE OF INVENTION: ANTI-MICROBIAL TARGETING CH
FILE REFERENCE: 2101363-991200
CURRENT APPLICATION NUMBER: US/10/077,624
CURRENT FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 09/910,358
PRIOR FILING DATE: 2001-07-19
PRIOR PRIOR PHIORIES DATE: 2001-07-19
PRIOR FILING DATE: 1909-08-20
NUMBER OF SEQ ID NOS: 31
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                                                 US-10-635-908-6 (1-119) x US-10-077-624-1 (1-563)
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                                                                                                                                                                                                                                                                                           LENGTH: 563
TYPE: DNA
ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                         FEATURE: OTHER INFORMATION: Synthesized using sequential PCR techniques
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1 AspValLysLeuValGluSerGlyGlyGlyLeuValLysLeuGlyGlySerLeuLysLeu
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Sequence 1, Application US/10706391

Sequence 1, Application US/10706391

Publication No. US20040137482A1

APPLICANT: THE REGENTS OF THE UNIVERSITY OF CALI

APPLICANT: WASHINGTON DENTAL SERVICE

APPLICANT: Gi, Fengxia

APPLICANT: APPLICATION NUMBER: US/10/706,391

CURRENT APPLICATION NUMBER: US 10/077,624

PRIOR APPLICATION NUMBER: US 10/077,624

PRIOR APPLICATION NUMBER: US 09/910,358

PRIOR APPLICATION NUMBER: US 09/910,358

PRIOR APPLICATION NUMBER: US 09/378,577

PRIOR APPLICATION NUMBER: US 09/378

PRIOR APPLICATION NUMBER: US 0
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; OTHER INFORMATION: Synthesized US-10-706-391-1
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US-10-706-391-1
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                                                         GGCTCCTACGGCTCCTATTACTATGCTATGGACTACTGGGGTCAAGGAACCTCAGTCACC 548
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Search completed: June 3, 2006, 07:32:36
Job time: 821.991 secs

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Title:
Perfect score:
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-MODEL-frame+ p2n.model -DEV=xlh
-MODEL-frame+ p2n.model -DEV=xlh
-Q-/abse/ABSSWEB_spool/US10635908/runat_02062006_104226_10433/app_query.fasta_1-Q-/abse/ABSSWEB_spool/US10635908/runat_0261062006_104226_10433/app_query.fasta_1-DB=Published_App_lications_NA_New -OFMT=fastap -SUFFIX=rnpbn -MINMATCH=0.1
-LOOPECL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -AMTRIX=blosum62
-TOOPECL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -FMATRIX=blosum62
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR_SCORE-pct -THR_MXX=100
-THR_MIN=0 -ALIGN=15 -MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXIEN=200000000 -HOST=abss05h
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Listing first 45 summaries
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-NO_MMAP -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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Copyright (c) 1993 - 2006 Biocceleration Ltd

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, App	equence 35,	equence 18, App	equence 15, App	equence 12, App	equence 57, App	puence 45,	equence 11, App	equence 17, App	equence 61, App	equence 167, Ap	equence 36	equence 19, App	equence 5, Appl	equence 41, App	equence 8, Appl	equence 173, Ap	e 14, App	equence 27, Apr	equence 65, App	nce 164, Ap	equence 20, App	equence 16, App	equence 65, App	equence 185, Ap	equence 1, Appl	equence 37, App	equence 90, App	equence 63, App	equence 129,	ence 69, App	equence 63, App	equence 53,	equence 95, App	equence 49, App	quenc	:

## ALIGNMENTS

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APPLICANT: BAILLY, VERONIQUE
APPLICANT: BAPLICANT: BAPLICANT: BROWNING, JEFFREY L.
APPLICANT: BROWNING, JEFFREY L.
TITLE OF INVENTION: MULTIVALENT LYMPHOTOXIN BETA RECEPTOR AGONISTS AND TITLE OF INVENTION: THERAPEUTIC USES THEREOF FILE REFERENCE: BUNALBECN
CURRENT APPLICATION NUMBER: US/11/155,444
CURRENT FILING DATE: 2005-06-17
PRIOR APPLICATION NUMBER: POT/US03/041393
PRIOR APPLICATION NUMBER: 60/435,154
PRIOR APPLICATION NUMBER: 60/435,154
PRIOR APPLICATION NUMBER: 60/435,154
PRIOR APPLICATION NUMBER: 60/435,155
PRIOR APPLICATION NUMBER: 60/435,185
PRIOR FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: 60/435,185
PRIOR FILING DATE: 2002-12-20
PRIOR FILING DATE: 2002-12-20
SOFTWARE: PATENTIN VEY: 3.3
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SOFTWARE: DNA
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US-10-635-908-6 (1-119) x US-11-155-444-1 (1-2094)
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                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: Heavy chain of OTHER INFORMATION: huCBEll/huBHAlO bispecific-1 antibody construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
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Sequence 7. Application US/1115444

Publication No. US20060104971A1

GENERAL INFORMATION:
APPLICANT: BARBER, ELLEN

APPLICANT: BAILLY, VERONIQUE

APPLICANT: BAILLY, VERONIQUE

APPLICANT: BROWNING, JEFFREY L.

TITLE OF INVENTION: THERAPEUTIC USES THEREOF

FILE REFERENCE: BGNA168CN

CURRENT APPLICATION NUMBER: US/11/155,444

CURRENT FILING DATE: 2005-06-17

PRIOR APPLICATION NUMBER: 60/135,154

PRIOR FILING DATE: 2003-12-20

PRIOR APPLICATION NUMBER: 60/435,154

PRIOR FILING DATE: 2003-12-20

PRIOR APPLICATION NUMBER: 60/435,185

PRIOR FILING DATE: 2002-12-20

PRIOR APPLICATION NUMBER: 60/435,185

PRIOR FILING DATE: 2002-12-20

PRIOR APPLICATION NUMBER: 60/435,185

PRIOR FILING DATE: 2002-12-20

PRIOR FILING DATE: 2002-12-20

NUMBER: OF SEQ ID NOS: 23

CONTRADED: THE PRIOR THE PRIOR THE PRIOR THE PRIOR FILING DATE: 2002-12-20

NUMBER: OF SEQ ID NOS: 23
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US-11-155-444-7
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Best Local Similarity:
Query Match:
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LENGTH: 2106
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                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: Heavy chain OTHER INFORMATION: huCBEll monospecific-1 antibody construct
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Conservative:
Mismatches:
Indels:
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NUMBER OF SEQ ID NOS: 23
SEQ ID NOS: 23
SEQ ID NO 5
EAUGH: 2196
TYPE: NO.
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Query Match:
DB:
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                                                                                                                                                                                                                                                                                                                                                                                                                 Alignment Scores:
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CURRENT APPLICATION NUMBER: US/11/155,444
CURRENT FILING DATE: 2005-06-17
PRIOR APPLICATION NUMBER: CT/US03/041393
PRIOR FILING DATE: 2003-12-27
PRIOR APPLICATION NUMBER: 60/435,154
PRIOR FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: 60/435,185
PRIOR APPLICATION NUMBER: 60/435,185
PRIOR FILING DATE: 2002-12-20
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Publication No. US20060104971A1

GENERAL INFORMATION:
APPLICANT: GARBER, ELLEN
APPLICANT: BAILLY, VERONIQUE
APPLICANT: BROWNING, JEFFREY L.
TITLE OF INVENTION: MULTIVALENT LYMPHOTOXIN BETA RECEPTOR AGONISTS.
TITLE OF INVENTION: THERMAPEUTIC USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
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LeuGlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArg 100
                                                   LeuAspThrValLysGlyArgPheThrIleSerArgAspAsnAlaLysAsnThrLeuTyr 80
                                                                                                                       ProGluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyr 60
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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: GARBER, ELLEN

APPLICANT: BAILLY, VERONIQUE

APPLICANT: BEROWNING, JEFFREY L.

TITLE OF INVENTION: MULTIVALENT LYMPHOTOXIN BETA RECEPTOR AGOI

TURNET FILING DATE: 2005-06-17

PRIOR APPLICATION NUMBER: 607435,154

PRIOR APPLICATION NUMBER: 607435,154

PRIOR APPLICATION NUMBER: 607435,185

PRIOR APPLICATION NUMBER: 607435,185

PRIOR APPLICATION NUMBER: 607435,185

PRIOR APPLICATION NUMBER: 607435,185

PRIOR FILING DATE: 2002-12-20

NUMBER OF SEQ ID NOS: 23

SOFTWARE: Patentin Ver: 3.3

SEQ ID NO 9

LENGTH: 2208

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE:

CHEEN TARGEMETER: DAGESTIALION OF Artificial Sequence: bucbe
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Best Local Similarity:
Query Match:
DB:
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US-11-155-444-9
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OTHER INFORMATION:
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Publication No. US20060104971A1
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                   101 SerGly---TyrPheSerMetAspTyrTrpGlyGlnGlyThrSerValThrValSerSer 119
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85.0%
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Matches:
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RESULT 6 US-11-154-103-27

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US-11-154-103-28

Sequence 28, Application US/11154103

PUBLICATION NO. US20060099205A1

GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: HORAK, EVA M.
APPLICANT: WEINER, LOUIS M.
APPLICANT: JAMES, MARKS D.
TITLE OF INVENTION: BISECCIFIC SINGLE CHAIN FV
FILE REFERENCE: 407T-000420US
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Best Local Similarity:
Query Match:
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NUMBER OF SEQ ID NOS: 57
SOFTWARE: PatentIn version 3.3
SEQ ID NO 27
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APPLICANT: HORAK, EVA M.
APPLICANT: WEINER, LOUIS M.
APPLICANT: JAMES, MARKS D.
TITLE OF INVENTION: BISPECIFIC SINGLE CHAIN FV ANTIBODY MOLECULES AND METHODS OF USE
TITLE OF INVENTION: THEREOF
FILE REFERENCE: 407T-000420US
CURRENT APPLICATION NUMBER: US/11/154,103
CURRENT FILING DATE: 2005-06-15
PRIOR APPLICATION NUMBER: US 60/370,276
PRIOR FILING DATE: 2002-04-02
PRIOR FILING DATE: 2002-04-02
PRIOR APPLICATION NUMBER: US 10/406,830
CURRENT APPLICATION NUMBER: US/11/154,103
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ORGANISM: Artificial
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Conservative:
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Indels:
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APPLICANT: HORAK, EVA M.

APPLICANT: HORAK, EVA M.

APPLICANT: WEINER, LOUIS M.

APPLICANT: JAMES, MARKS D.

ITILE OF INVENTION: BISERCIFIC SINGLE CHAIN FV ANTIBODY MOLECULES AND METHODS OF USE

TITLE OF INVENTION: THEREOF

FILE REFERENCE: 407T-000420US

CURRENT FILING DATE: 2005-06-15

PRIOR APPLICATION NUMBER: US/11/154,103

CURRENT FILING DATE: 2005-06-15

PRIOR APPLICATION NUMBER: US 60/370,276

PRIOR APPLICATION NUMBER: US 60/370,276

PRIOR FILING DATE: 2003-04-02

PRIOR FILING DATE: 2003-04-04

NUMBER OF SEQ ID NOS: 57

SOFTWARE: Patentin version 3.3

SEQ ID NO 30

LENGTH: B70

TYPE: DNA

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PRIOR APPLICATION NUMBER: US 60/370,276;
PRIOR FILING DATE: 2002-04-02;
PRIOR APPLICATION NUMBER: US10/406,830;
PRIOR FILING DATE: 2003-04-04;
NUMBER OF SEQ ID NOS: 57;
SOFTWARE: Patentin version 3.3;
SEQ ID NO 28;
LENGTH: 873;
TYPE: DNA
ORGANISM: Artificial
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US-11-154-103-30
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                                                                                                                                                                                                                                                                                                                                                                           Sequence 30, Application US/111 Publication No. US20060099205A1 GENERAL INFORMATION:
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21

189

249

309

USE

ORGANISM: Artificial

Percent Similarity:
Best Local Similarity:
Query Match:
DB:

2.11e-49 437.50 85.7% 73.1% 70.9%

Length: Matches:

Mismatches: Indels: Conservative:

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; FEATURE: ; OTHER INFORMATION: US-11-154-103-30

Nucleic acid encoding single

chain antibody

Alignment Scores

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                                                ; OTHER INFORMATION: ; OTHER INFORMATION: US-11-216-033-3
Alignment Scores: Pred. No.:
                                                                                                                                                                                                                    APPLICANT: QU, ZHENGXING
TITLE OF INVENTION: PRODUCTION AND USE OF NOVEL PEPTIDE-BASED AGENTS |
TITLE OF INVENTION: USE WITH BI-SPECIFIC ANTIBODIES
FILE REFERENCE: 40923-0074US4
CURRENT APPLICATION NUMBER: US/11/216,033
CURRENT FILING DATE: 2005-09-01
PRIOR APPLICATION NUMBER: US/09/823,746
PRIOR FILING DATE: 2001-04-03
PRIOR APPLICATION NUMBER: 09/337,756
PRIOR APPLICATION NUMBER: 09/337,756
PRIOR FILING DATE: 1999-06-22
NUMBER OF SEO ID NOS: 18
SOPTWARE: Patentin Ver. 3.3
LENGTH. 354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3, Application US/11216033
Publication No. US20060104899A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: HANSEN, HANS J.
APPLICANT: GRIFFITHS, GARY L.
APPLICANT: MCBRIDE, WILLIAM J.
APPLICANT: LEUNG, SHUI-ON
                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                      NAME/KEY: CDS
LOCATION: (1)..(354)
                                                                                                                                                         FEATURE:
                                                                                                                                                                      TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                       FEATURE:
                                                                                                                                                                                                           ENGTH: 354
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -GlyTyrPheSerMetAspTyrTrpGlyGlnGlyThrSerValThrValSerSer 119
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                                                                  Description of Artificial Sequence: Synthetic nucleotide sequence
Length
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:

Conservative: Mismatches: Indels:

Score:

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APPLICANT: ROUTLEDGE, EDWARD G.
APPLICANT: WALMANN, HERMAN
TITLE OF INVENTION: HUMANIZED ANTI-CD3 SPECIFIC ANTIBODIES (AS AMENDED)
FILE REFERENCE: 604-704
CURRENT FILING DATE: 2005-12-15
PRIOR APPLICATION NUMBER: 10/743.423
PRIOR APPLICATION NUMBER: 10/743.423
PRIOR FILING DATE: 203-12-23
PRIOR FILING DATE: 1995-06-07
PRIOR APPLICATION NUMBER: 08/478,684
PRIOR APPLICATION NUMBER: 09/478,684
PRIOR APPLICATION NUMBER: 0906-07
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Best Local Similarity:
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                                                                                                                                                                   OTHER INFORMATION: Description OTHER INFORMATION: sequence US-11-300-563-23
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                                                                                                        Pred. No.:
                                                                                                                          Alignment Scores:
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                                                                                                                                                                                                                                                                                                SEQ ID NO 23
LENGTH: 357
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Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: BOLT,
APPLICANT: CLAR
APPLICANT: GORM
                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 26
                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                              TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               121 GAAAAGAGGCTGGAGTGGGTCGCAACCCTGAGTGGTGATGGTGATGACATCTACTATCCA 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               82 GlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArgSer 101
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4.47e-49
431.00
84.9%
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69.9%
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Conservative:
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Publication No. US/20060093600A1
GENERAL INFORMATION:
APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, ROMALD P.
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
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US-11-211-917-49
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FILE REPERENCE: ABX-PP/3 US
CURRENT APPLICATION NUMBER: US/11/211,917
CURRENT FILLING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR FILING DATE: 2002-11-08
PRIOR APPLICATION NUMBER: 60/348,980
PRIOR FILING DATE: 2001-11-09
PRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
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DB:
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Pred. No.:
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SEQ ID NO 49
LENGTH: 373
TYPE: DNA
ORGANISM: Homo sapiens
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GluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyrLeu
                                                                                          CysAlaAlaSerGlyPheThrPheSerAsnTyrTyrMetSerTrpValArgGlnThrPro
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                                                                                                                                        GTGCAGCTGGTGGAGTCTGGGGGAGGCGTGGTCCAGCCTGGGAGGTCCCTGAGACTCTCC
                                                                      TGTGCAGCCTCTGGATTCACCTTCAGTCGCTATGGCATGCACTGGGTCCGCCAGGCTCCA
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429.50
80.5%
69.1%
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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GENERAL INFORMATION:

APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONALD P.
APPLICANT: GLADUE, RONALD P.
APPLICANT: GLADUE, APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
TITLE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/11/211,917
CURRENT FILING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR FILING DATE: 2001-11-09
PRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
NUMBER OF SEQ ID NOS: 147
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Best Local Similarity:
Query Match:
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US-11-211-917-95
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Alignment Scores:
Pred. No.:
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Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 373
TYPE: DNA
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                       82 GlnMetSerSerLeuLysSerGluAspThrAlaLeuPheTyrCysAlaArgHisArgSer 101
                                                                               42 GluLysArgLeuGluLeuValAlaAlaIleAsnSerAspGlyGlyIleThrTyrTyrLeu
                                                                                                                                                                                         64 TGTGCAGCCTCTGGATTCACCTTCAGTCGCTATGGCATGCACTGGGTCCGCCAGGCTCCA
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                                                             GACTCCGTGAAGGGCCGATTCACCATCTCCAGAGACAATTCCAAGAACACGCTGTATCTG
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No. US20060093600A1
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TITLE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/11/211,917
CURRENT FILING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR FILING DATE: 2002-11-08
PRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: PACENTIN VET. 2.1
SEQ ID NO 53
LENGTH: 1410
TYPE: DNA
ORGANISM: Homo sapiens
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Publication No. US20060093600A1
GENERAL INFORMATION:
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APPLICANT: GLADUE, RONALD P.
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
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                                                       GGAAAGACTTACTACCACTACTGTGGTATGGACGTCTGGGGCCAAGGGACCACGGTCACC
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GTCTCCTCA 429
                          ValSerSer 119
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RESULT 14

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SOFTWARE: Patentin version 3.1
SEQ ID NO 63
LENGTH: 369
TYPE: DNA
CORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (1)..(369)
OTHER INFORMATION:
US-10-546-594-63
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Publication No. US20060088538A1
GENERAL INFORMATION:
APPLICANT: HOSOKOWAM, Saiko
APPLICANT: HOSOKOWAM, Saiko
APPLICANT: AOKI, Masahiko
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APPLICANT: FUKUDA, KAZUMBAB
IITLE OF INVENTION: MONOCLONAL ANTIBODY AND GENE ENCODING THE SAME, HYBRIDOMA, PHARW
IITLE OF INVENTION: COMPOSITION, AND DIAGNOSTIC REAGENT
FILE REFERENCE: 238067
CURRENT APPLICATION NUMBER: US/10/546,594
CURRENT FILING DATE: 2005-08-19
PRIOR APPLICATION NUMBER: JP 2003/54670
PRIOR PILING DATE: 2003-02-28
PRIOR PILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 132
CONTACTOR OF SEC ID NOS: 132
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UMEKI, Hiroe
SAIKAWA, Yoshiro
KUMAI, Koichiro
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LENGTH: 369
LENGTH: 369
TYPE: DNA
ORGANISM: Homo sapiens
PEATURE:
NAME/KEY: CDS
LOCATION: (1)..(369)
OTHER INFORMATION:
US-10-546-594-69
US-10-546-594-69
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Publication No. US20060088538A1
GENERAL INFORMATION:
APPLICANT: HOSOKAWA, Saiko
APPLICANT: HOSOKAWA, Saiko
APPLICANT: HOSOKAWA, YOA
APPLICANT: HITAKAWA, YOA
APPLICANT: ITAMI, Seima
APPLICANT: UMEKI, Hiroe
APPLICANT: UMEKI, Hiroe
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Best Local Similarity:
Search completed: June 3, 2006, 09:17:13 Job time: 16.5796 secs
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PRIOR FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: JP 2003/194643
PRIOR FILING DATE: 2003-07-09
NUMBER OF SEQ ID NOS: 132
SOFTWARE: PatentIn version 3.1
SEQ ID NO 69
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APPLICANT: UMEKI, Hiroe
APPLICANT: SAIKAWA, Yoshiro
APPLICANT: KUMAI, Koichiro
APPLICANT: FUKUDA, Kazumasa
TITLE OF INVENTION: MONOCLONAL ANTIBODY AND GENE ENCODING THE SAME, HYBRIDOMA, PHARMA
TITLE OF INVENTION: COMPOSITION, AND DIAGNOSTIC REAGENT
FILE REFERENCE: 238067
CURRENT APPLICATION NUMBER: US/10/546,594
CURRENT FILING DATE: 2005-08-19
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                                                                                                                                      101 SerGlyTyr------PheSerMetAspTyrTrpGlyGlnGlyThrSerValThr 116
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-MODEL=frame+ p2n.model -DEV=xlh
-Q=/abs=/ABSSWEB spool/US10635908/runat 02062006_104214_10235/app_query.fasta_1
-Q=/abs=/ABSSWEB spool/US10635908/runat 02062006_1-1-LOOPCL=0 -LOOPEXT=0
-DB=CenEmbl -QFMT=fastap -SUFFIX=zge -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
-DOCALIGN=200 -THR SCORE=pct -THR MAX=100 -THR MIN=0 -ALIGN=15 -MODE=LOCAL
-OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=20000000000 -HOST=abss05h
-USER-US10635908 @CGN 1 1 5767 @xunat 02062006 104214 10225 -NCPU=6 -ICPU=3
-NO MMAP -NEG SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG -DEV_TIMEOUT=120
-NARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                               Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                               Score
1030
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Xgapop 10.0 , Xgapext
Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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## ALIGNMENTS

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Gonzalez,T.N., Leong,S.R. and Presta,L.G.
Humanized anti-IL-8 monoclonal antibodies
Patent: US 6133426-A 24 17-CCT-2000;
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Zapata, G.A.
Antibody fragment-polymer conjugates and monoclonal antibodies
Patent: JP 2001512489-A 25 21-AUG-2001;
GENENTECH INC
                                                                                        BD069191
Antibody fragment-polymer monoclonal antibodies.
BD069191
                                    Hsei, V., Koumenis, I., Leong, S.R., Presta, L.G.,
                                          unclassified sequences.
1 (bases 1 to 714)
                                                           unidentified
                                                                  unidentified
                                                                         BD069191.1 GI:22614794
JP 2001512489-A/25.
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pN JP 2001512499-A/25
pD 21-AUG-2001
PF 2001512499-A/25
pP 21-FB-1998 JP 1998536905
pF 20-FBB-1998 US 09/012116 PI
PR 21-FBB-1997 US 08/804444,22-JAN-1998 US 09/012116 PI
PR 21-FBB-1997 US 08/804444,22-JAN-1998 US 09/012116 PI
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LeuSerLysalaAspTyrGluLysHisLysValTyrAlaCysGluValThrHisGlnGly
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C12N15/13, C07K19/00, A61K47/48, C07K16/24, C12N15/85, C12N5/10
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Antibody fragment-polymer conjugates and monoclonal antibodies
Patent: EP 1415998-A 24 06-MAY-2004;
Genentech, Inc. (US)
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other sequences; artificial sequences.
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Sequence 24 from Patent
CQ802834
CQ802834.1 GI:47109852
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GluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThr
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                                                                                                                                                GGGACCAAGCTGGAGCTTCGAAGAGCTGTGGCTGCACCATCTGTCTTCATCTTCCCGCCA
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                                                                                             TCTGATGAGCAGTTGAAATCTGGAACTGCTTCTGTTGTGTGCCTGCTGAATAACTTCTAT
                                          CCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAG
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Zapata, G.

Methods of treating inflammatory disease
fragment-polymer conjugates
patent: US 6458355-A 24 01-OCT-2002;
patenti US 6458355-A 25 01-OCT-2002;
Genentech, Inc.; South San Francisco, CA
Location/Qualifiers
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ProArgGluAlaLysValGlnTrpLysValAspAsnAlaLeuGlnSerGlyAsnSerGln
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                                                                                                                                                                    CGCTTCACAGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCCATGTGCAGTCT
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                                 TCTGATGAGCAGTTGAAATCTGGAACTGCTTCTGTTGTGTGCCTGCTGAATAACTTCTAT
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/mol_type="genomic |
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Methods of treating inflammatory diseases
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patent: US 6468532-A 24 22-OCT-2002;
Genentech, Inc.; South San Francisco,
Location/Qualifiers
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SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr
                                                                           ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer
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                                                                                                                      GGGACCAAGCTGGAGCTTCGAAGAGCTGTGGCTGCACCATCTGTCTTCATCTTCCCGCCA
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/mol_type="genomic DNA"
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Sequence
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Fong, S., Hebert, C.Alice., Kim, K.Jin. Anti-IL-8 antibody fragments
Patent: US 5677426-A 27 14-OCT-1997;
Location/Qualifiers
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                                                                                                                             ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer
                                                                                                                                                          GGGCAATCTCCTAAAGCACTGATTTACTCGTCATCCTACCGGTACAGTGGAGTCCCTGAT
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  SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr 140
                                     GlyThrLysLeuGluIleLysArgThrValAlaAlaProSerValPheIlePheProPro 120
                                                                                                                GAAGACTTGGCAGACTATTTCTGTCAGCAATATAACATCTATCCTCTCACGTTCGGTCCT
                         GGGACCAAGCTGGAGCTTCGAAGAGCTGTGGCTGCACCATCTGTCTTCATCTTCCCGCCA
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from patent US 5677426.
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Doerschuk, C.M., Fong, S., Hebert, C.Alice.,
Methods for treating bacterial pneumonia
Patent: US 5686070-A 27 11-NOV-1997;
Location/Qualifiers
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1 (bases 1 to 714)
Fong, S., Hebert, C.Alice., Kim, K.Jin. and Methods for treating ulcerative colltise Patent: US 5707622-A 27 13-JAN-1998;
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GlyThrLysLeuGluIleLysArgThrValAlaAlaProSerValPheIlePheProPro|
                                                       CGCTTCACAGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCAGCCATGTGCAGTCT
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1 (bases 1 to 714)

1 (bases 1 to 714)

Doerschuk,C.M., Fong,S., Hebert,C.Alice., Kim,K.Jin. and Leong,S. Anti-IL-8 monoclonal antibodies for treatment of inflammatory disorders

Patent: US 5702946-A 27 30-DEC-1997;
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                                                    ArgPheThrG1ySerG1ySerG1yThrAspPheThrLeuThrI1eSerAsnMetG1nSer
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/mol_type="unassigned DNA"
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AR647560
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Hsei V. Koumenis, I., Leong, S.R., Presta, L.G., Shahrokh, Z. Zapata, G.A.
Antibody fragment-polymer conjugates and humanized anti-IL-
monoclonal antibodies
patent: US 6870033-A 24 22-MAR-2005;
Genentech, Inc.; South San Francisco, CA
Location/Qualifiers
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                            Patent: WO 2005108989-A 10 17-NOV-2005; GENENTECH, INC. (US)
                                                                                                                                                                                                                                                                                                                              Chuntharapai, A
                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                   10 from Patent
                                                                                                                                                                                                                                               /organism="Mus musculus"
/mol type="unassigned DN/
/db xref="taxon:10090"
1684
                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                      /note="Unknown amino acid
                                                                                                                                                     1.06e-111
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                                                                                                     Percent Similarity:
Best Local Similarity:
                                                                                                                                Score:
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                                                                                                                                                                                                                                    hematopoietic origin
Patent: WO 2005063299-A 14 14-JUL-2005;
Genentech, Inc. (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                      Chang, W.
                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                        Compositions and methods
                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (human)
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      GATATCGTGATGACCCAGTCTCATAAATTCATGTCCACATCAGTAGGAGACAGGGTCAGC
                   AspIleValMetThrGlnSerGlnArgPheMetSerThrThrValGlyAspArgValSer
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/mol_type="unassigned DN
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beta lactamase; bla gene; Fab fragment
beta lactamase; blafene; Fab fragment
IGK gene; light chain; tet repressor;
Cloning vector pASK88-huHRS3-VH-EP3/1
Cloning vector pASK88-huHRS3-VH-EP3/1
                                                                                                                                                                                                                   Submitted (22-OCT-2004) Schlapschy M., LS Biologische Muenchen, An der Saatzucht 5, 85354 Freising, GERMANY Location/Qualifiers
1. 4604
                                                                                                                                                                                                                                                                                                                                                                                         numentization of an anti-CD30 Fab Fragment as a reagent for the immunotherapy of Hodgkin's Lymphoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AJ852004
Synthetic construct
Fab fragment.
                                                                                                                                                                                                                                                                                                                                       Schlapschy, M.
                                                                                                                                                                                                                                                                                                                                                                               Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCysGluValThrHisGlnGly 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThr 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TCTGATGAGCAGTTGAAATCTGGAACTGCTTCTGTTGTGTGCCTGCTGAATAACTTCTAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CGCTTCACTGGCAGTGGATCTGGGACGGATTTCACTTTCACCATCAGCAGTGTGCAGGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GGACATTCTCCTAAACTACTGATTTACTCG---GGATACCGGTACACTAGAGTCCCTGAT
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                                                                                                                                                                                                                                                                                                                                                            (bases 1 to 4604)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequences; artificial sequences;
                                                                                                                                                                                                                                                                                                                    Submission
/note="tetA promoter region"
45. .50
69. .74
95. .97
                                                                                                                                                     /organism="Cloning vector pASK88-huHRS3-VH-EP3/1"
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                                                                                                          /rearranged
/lab_host="
                                                                                                             host="Escherichia coli"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4604 bp DNA
pASK88-huHRS3-VH-EP3/1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      circular SYN 08-JAN-2005 for humanized anti-CD30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene.
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3180. .3803
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2310. .3170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /codon_start=1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /product="light chain"
1647. .1721
/note="lipoprotein transcription terminator"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HQGLSSPVTKSFNRGEC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                210. .905
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/note="ompA signal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /codon_start=1
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                                                                                                                                                                                      EIGASLIKHW"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="intergenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          991. .1632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /gene="IGK"
/note="phoA signal peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /gene="IGHG1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /gene="IGK"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'note="mouse-human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'gene="IGK"
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Score:
Percent Similarity:
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                                                                              GAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTCAGCAGCACCCTGACG
                                                                                                                      GluAspLeuAlaAspPhePheCysGlnGlnTyrSerAsnTyrProTrpThrPheGlyGly 100
                                                                                                                                                                                                                                                                                                   ArgPheThrG1ySerG1ySerG1yThrAspPheThrLeuThrI1eSerAsnMetG1nSer
          LeuSerSerProValThrLysSerPheAsnArgGlyGluCys 214
                                       CTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGC
                                                                                           GluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThr
                                                                                                                                                               TCTGATGAGCAGTTGAAATCTGGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTAT
                                                                                                                                                                           SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr
                                                                                                                                                                                                       GGCACCAAACTCGAGATCAAACGGACTGTGGCTGCACCATCTGTCTTCATCTTCCCGCCA 1350
                                                                                                                                                                                                                    GlyThrLysLeuGluIleLysArgThrValAlaAlaAroSerValPheIlePheProPro
                                                                                                                                                                                                                                                GAAGACTTCGCTACCTACTGCCAGCAGTACCACACCTACCGGTTAACCTTCGGTCAG
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3930. 4603
/note="Colei ori"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        direction=RIGHT
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962.00
92.5%
86.0%
86.2%
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Matches:
Conservative:
Mismatches:
Indels:
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184
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Search completed: June 3, 2006, 06:56:30 Job time: 6313.61 secs

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Command line parameters:

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-DB=N Geneseq -QFMT=fastap -SUFFIX=rng -MINMATCH=0.1 -LÖOPCL=0 -LÖOPEXT=0
-UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
-DOCALIGN=200 -THR SCORE-pct -THR MAX=100 -THR MIN=0 -ALIGN=15 -MODE=LOCAL
-OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=2000000000 -HOST=abss05h
-USER=US10635908 @CGN 1 1 942 @runat 02062006 104212 10223 -NCPU=6 -LCPU=3
-USER=US10635908 @CGN 1 1 942 @runat 02062006 104212 10223 -NCPU=6 -LCPU=3
-NO MMAP -NEG SCÖRES=0 -WAIT -DSPBLOCK=100 -LÖNGLOG -DEV_TIMEOUT=120
-WARN TIMEOUT=30 -THREADS=1 -XCAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
-YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                           Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Sequence:
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                                                                                                                                                                        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                           Score
1030
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seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N_Geneseq_8:*
1: geneseqn19
2: geneseqn20
3: geneseqn20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human C5E10 antibody light chain DNA.
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                                                           29-JAN-2002; 2002WO-US002373
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/product= "Human C5E10 light chain protein"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Domain deleted CC49 antibodies comprise a heavy chain human CC49 domain deleted sequence in which CH2 domain has been deleted and are reactive with CD20 and comprise a heavy chain having a sequence of a derived domain deleted C2B8 construct where the CH2 domain has been deleted. Sequences of the invention are useful for imaging a neoplasm. They are also useful for treating myelosuppressed patients suffering from neoplastic disorder such as haematologic neoplasm, preferably non-complastic disorder, colon cancer and haematologic malignancy. They are useful for reducing tumour size, inhibiting tumour growth and/or prolonging the survival time of tumour-bearing animals and for treating tumours. The present sequence is a DNA encoding human C5E10 light chain protein. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel domain deleted CC49 antibody reactive with tumor associated -72, or C2B8 antibody reactive with CD20, useful for treating myelosuppressed patient suffering from a neoplastic disorder.
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                                                                                           SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr
                                                                                                                                                                                                      GlyThrLysLeuGluIleLysArgThrValAlaAlaProSerValPheIlePheProPro
                                                                                                                                                                                                                                                                                      GAAGACTTGGCAGATTATTTCTGTCAGTTATATAGCAGCTATCCTCTCACGTTCGGAGGG
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ProArgGluAlaLysValGlnTrpLysValAspAsnAlaLeuGlnSerGlyAsnSerGln
                                                                                                                                                                        GGGACCAAGCTGGAAATCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTTCCCGCCA
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         monomeric subunits, where the monomeric subunits are non-covalently associated. (I) is useful for treating a disorder, especially immune disorder, and neoplastic disorder such as relapsed Hodgkin's disease, resistant Hodgkin's disease high grade, low grade and intermediate grade non-Hodgkin's lymphomas, B cell chronic lymphocytic leukemia (B-CLL), lymphopalasmacytoid lymphoma (FD), mantle cell lymphoma (MCL), follicular lymphoma (FL), diffuse large cell lymphoma (DLCL), Burkitt's lymphoma, AIDS-related lymphomas, monocytic B cell lymphoma, angioinmunoblastic lymphodenopathy, small lymphocytic, follicular, diffuse large cell, diffuse small cleaved cell, large cell immunoblastic lymphoblastoma, small, non-cleaved, Burkitt's and non-Burkitt's, follicular, mixed small
                                                                                                                                                                                                                                                                                                                                                                                                          29-JAN-2001; 2001US-0264318P.
16-NOV-2001; 2001US-0331481P.
21-DEC-2001; 2001US-0341858P.
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                                                                                                                                                                                                                                                    Novel dimeric antibody useful for treating immune disorder and neoplastic disorder, has several non-covalently associated monomeric subunits.
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                                                                                                                                                                                         The invention relates to a dimeric antibody (I) comprising several
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DB; ABB82839.
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           Xenotransplantation; graft rejection; cell interaction; vascular cell adhesion molecule; VCAM; monoclonal antibochimeric antibody; diagnosis; ss.
                                                                                   16-JUN-1997
                                                            Murine anti-porcine VCAM 2A2 light chain DNA sequence
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                                                                                                                                                                                 LeuSerSerProValThrLysSerPheAsnArgGlyGluCys
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                                                                                                                                   standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA sequences (AAT62929-30) respectively code for the light chain (AAW14931) and heavy chain (AAW14932) of murine anti-porcine soluble vascular cell adhesion molecule (VCAM) monoclonal antibody (MAb) 2A2. Hybridoma 2A2 was produced by standard techniques using recombinant, soluble porcine VCAM as immunogen. Chimeric antibodies can be produced by cloning MAb 2A2 and 3F4 (see also AAT62934-35) variable regions into expression plasmid pAPEX-3P modified to contain the human gamma4 constant region in place of the human gamma1 C1 region. Sequences are provided for 2A2 (chimeric) human G2/G4 cDNA (AAT62931), a 2A2 human G2/G4 expression plasmid insert (AAT62932), and a 2A2 human IGG4 expression plasmid insert (AAT62932), and a 2A2 human IGG4 expression plasmid insert (AAT62932), and a 2A2 human IGG4 expression plasmid insert (AAT62932). The chimeric antibodies are specific for poroline VCAM. They are useful for diagnosing human rejection of porcine xenotransplants and for improving xenotransplantation of porcine cells, tissues and organs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antibodies binding to porcine but not human cell interaction puuseful to treat and assay for rejection of xenografted porcine tissues or cells.
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P-PSDB; AAW14931.
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rcrgargagcagrigaaarcrggaacrgccrcrgrrgrgrgccrgcrgaaraacrrcrar SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr 4773

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GluAspLeuAlaAspPhePheCysGlnGlnTyrSerAsnTyrProTrpThrPheGlyGly

CGCTTCACAGGCAGTGGATCTGGGACAGATTTCACTCTCACCATCACCAATGTGCAGTCT ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer GGCCAGTCTCCTAAAACACTTATTTACTCGGCATCCTTCCGCTACAGTGGAGTCCCTGAT

GlyThrLysLeuGluIleLysArgThrValAlaAlaProSerValPheIlePheProPro GAAGACTTGGCAGAGTATTTCTGTCATCAATATAACTCCTATCCTCTCACGTTCGGGGGG

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Pred. No.:
                                                                                       US-10-635-908-15 (1-214) x AAT62933
                                                                                                                                                                                                                                                                                                                                          A DNA sequence (AAT62933) comprises a 2A2 human IgC4 expression plasmid insert sequence. It is obtd. by cloning the light chain (see also AAT62929) and heavy chain (see also AAT62930) variable region sequences of murine anti-porcine vascular cell adhesion molecule (VCAM) monoclonal antibody 2A2 into an expression plasmid modified to contain the human gamma4 constant region in plasmid of the human gamma C1 region. The encoded heavy chain (AAM14935) and light chain (AAM14935) can be used to produce a chimeric antibody that is specific for porcine VCAM. The chimeric antibody is useful for diagnosing human rejection of porcine xenotransplants and for improving xenotransplantation of porcine cells, tissues and organs into human recipients. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antibodies binding to porcine but not human cell interaction proteins - useful to treat and assay for rejection of xenografted porcine organs, tissues or cells.
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P-PSDB; AAW14935, AA
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26-SEP-1996;
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A CDNA insert (AAT03363) in vector pA51214VL' codes for the light chain variable region of anti-human interleukin-8 mouse monoclonal antibody 5.12.14 and human IgG1 light chain constant region. The cDNA was obtd. 1 amplification of cDNA derived from 5.12.14-producing hybridoma cells using primers (AAT03352-55) based on light chain sequences. It is used
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Interleukin-8; IL-8; antagonist; anti-interleukin-8 antibody; asthma; inflammatory disorder; neutrophil chemotaxis; elastase release;

monoclonal

antibody light chain region

encoding

CDNA

80

249 8

309

369

120

160

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489

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                                                                                                US-10-635-908-15 (1-214) x AAT78570
                                                                                                                                                                                                                                              Methods for treating asthma in mammals with interleukin-8 (II-8) antagonists (anti-II-8 antibodies) have been produced. Murine 5.12.14 monoclonal antibody is an anti-II-8 antibody. The present sequence encodes the murine 5.12.14 variable light chain region and the human IgG1 constant light chain region chimeric monoclonal antibody. Anti-II-8 antibodies are especially used to treat or prevent allergic asthma in humans. They inhibit: (a) neutrophil chemotaxis in response to II-8; (b) II-8-mediated release of elastase from neutrophils and (c) binding of II-8 to neutrophils. Anti-II-8 antibodies can be used to treat many other inflammatory disorders, e.g. ischaemic reperfusion, ARDS, dermatitis, particularly bacterial pneumonia and inflammatory bowel disease. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treating asthma by admin. that inhibits binding of ]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hebert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-JUN-1996;
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DB; AAW23786.
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                                              ABPIleValMetThrGlnSerGlnArgPheMetSerThrThrValGlyAspArgValSer
             IleThrCysLysAlaSerGlnAsnValValSerAlaValAlaTrpTyrGlnGlnLysPro
                                                                                                                                                                                                                        714 BP;
GTCACCTGCAAGGCCAGTCAGAATGTGGGGTACTAATGTAGCCTGGTATCAACAGAAACCA
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/label= Murine_5.12.14_variable_light_chain_region
397. .714
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   of interleukin-8 antagonist - pref. an antibody IL-8 to neutrophil(s), neutrophil chemotaxis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Chimeric_monoclonal_antibody_light_chain"
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                                                                                                (1-714)
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Conservative:
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                                                                                                                                                                                                                                                             murine light chain; 5.12.14; human interleukin-8; IL-8; anti IL-8; monoclonal antibody; diagnosis; inflammatory disorder; treatment; pneumonia; complementarity determining region; CDR; variable region; constant region; chimeric; pantiIL-8.2; netrophil chemotaxis; ds.
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24-FEB-1998
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Homo sapiens.
                                                                                                                                                                                                                                                                                                                             Chimeric monoclonal antibody
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGGCAATCTCCTAAAGCACTGATTTACTCGTCATCCTACCGGTACAGTGGAGTCCCTGAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GlyGlnSerProLysLeuLeuIleTyrSerAlaSerAsnArgTyrThrGlyValProAsp
                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LeuSerSerProValThrLysSerPheAsnArgGlyGluCys 214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCysGluValThrHisGlnGly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ProArgGluAlaLysValGlnTrpLysValAspAsnAlaLeuGlnSerGlyAsnSerGln
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                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                                                                                           /*tag= a
1. .69
1. .69
/*tag= b
/note= "signal |
70. .711
                                                          /note= "amino acids 24-132 are of murine variable light region and amino acids 133-237 are of human constant light region "
                                                                                                                                                                                      Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 714 BP;
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                                                                                                                                                                GluAspLeuAlaAspPhePheCysGlnGlnTyrSerAsnTyrProTrpThrPheGlyGly 100
                                                                                                                                                                                                                                                                                                     GTCACCTGCAAGGCCAGTCAGAATGTGGGTACTAATGTAGCCTGGTATCAACAGAAACCA
ProArgGluAlaLysValGlnTrpLysValAspAsnAlaLeuGlnSerGlyAsnSerGln 160
                                                               SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr 140
                                                                                                               GlyThrLysLeuGluIleLysArgThrValAlaAlaProSerValPheIlePheProPro 120
                                                                                                                                                    GAAGACTTGGCAGACTATTTCTGTCAGCAATATAACATCTATCCTCTCACGTTCGGTCCT 369
                                                                                                                                                                                                     ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer
                                                                                                   GGGACCAAGCTGGAGCTTCGAAGAGCTGTGGCTGCACCATCTGTCTTCATCTTCCCGCCA
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Matches:
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RESULT 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammtory disorder; bacterial pneumonia; neutrophil chemotaxis; anti IL-8 monoclonal antibody; IL-8 mediated elastase release; streptococcus pneumoniae; Escherichia coli; Pseudomonas aeruginos
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15-APR-1998
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inflammtory o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Murine variable and human IgG1 constant region light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAR-1995;
                                                                                                                                               Treatment of bacterial pneumonia - with monoclonal antibody interleukin-8; inhibits lung inflammatory conditions.
                                                                                                                                                                                                                  WPI; 1997-558085/51.
P-PSDB; AAW42319.
                                                                                                                                                                                                                                                                                Kim KJ,
                                                                                                                                                                                                                                                                                                                                                                                       03-MAR-1994;
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GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
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/*tag=
396. .7
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/note= "encodes the variable light chain region
/nurine antibody 5.12.14"
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1. .395
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1. .714
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IgG1"
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The present sequence depicts the coding sequence for murine monoclonal antibody 5.12.14 light chain variable region and the human IgG1 light chain constant region. The 5.12.14 antibody is a murine anti interleuk: 8 (IL-8) IgG2a isotype. IL-8 is a neutrophil chemotactic peptide secret

Fig 19; 63pp; English.

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Pred. No.:
Score:
RESULT 9
AAV06413
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   by a variety of cells in response to inflammatory mediators. IL-8 can play an important role in the pathogenesis of inflamamory disorders such as adult respiratory distress syndrome (ARDS), septic shock and multiple organ failure. Treatment of bacterial pneumonia in a mammal comprises administering an anti IL-8 monoclonal antibody, such as, e.g humanised 5.12.14, that binds human IL-8 with a Kd of 10-8 to 10-11 M, inhibits neutrophil chemotaxis in response to IL-8, inhibits IL-8 mediated elastase release by neutrophils and does not bind to C5a, beta-TG or platelet factor 4. IL-8 specific monoclonal antibodies are especially for treating pneumonia caused by Streptococcus pneumoniae, E. coli or Pseudomonas aeruginosa in humans. The antibodies may also be used in the treatment of ulcerative colitis and other inflammatory conditions. (Updated on 17-OCT-2003 to standardise OS field)
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                                                                                  LeuSerSerProValThrLysSerPheAsnArgGlyGluCys 214
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                                                                                                                                                LeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCysGluValThrHisGlnGly
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AAV06413

standard; DNA;

ВP

US-10-635-908-15 (1-214) x AAV06413

(1-714)

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Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                                                                                            This DNA encodes for the chimeric Fab 5.12.14 which comprises of complementarity determining regions (CDRs) of a light chain variable region of the murine species and light chain constant region of the human species and light chain constant region of the human species and light chain constant region of the human species. This chimeric Fab is used to construct a plasmid pantiII-8.2 which consists of murine-human variable/constant regions cooth the light and heavy chains of 5.12.14 yearious standard recombinant techniques. The recombinant human-murine chimeric Fab has anti interleukin-8 (II-8) activity and can inhibit the II-8 binding to interleukin-8 mediated inflammatory diseases and in the treatment of ulcerative colitis and bacterial pneumonia. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        murine light chain; 5.12.14; human interleukin-8; IL-8; anti IL-8 monoclonal antibody; diagnosis; inflammatory disorder; treatment; pneumonia; complementarity determining region; CDR; variable regionstant region; chimeric; pantiIL-8.2; ulcerative cloitis; ds.
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                                                                                                                                                          Sequence 714
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Monoclonal antibody specific for interleukin-8 - used for treating mediated inflammatory diseases, e.g. ulcerative colitis.
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1. .69
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/*tag= b
/note= "signal peptide of STII"
70. .711
70. .711
/*tag= c
/note= "amino acids 24-132 are of murine variable light region and amino acids 133-237 are of human constant light region "
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1.55e-81
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                                                                                                                                                                                                  Humanised antibody; chimeric antibody; monoclonal antibody; mouse; human; Fab; interleukin-8; inflammation; immunotherapy; psoriasis; inflammatory bowel disease; Crohn's disease; ulcerative colitis; ischaemic reperfusion; adult respiratory distress syndrome; dermatitis; meningitis; encephalitis; uveitis; autoimmune disease; rheumatoid arthritis; Sjorgen's syndrome; vasculltis; leukocyte diapedesis; multiple organ injury syndrome; septicaemia; urauma; alcoholic hepatitis; pneumonia; pleurisy; alveolitis; vasculitis; bronchiectasis; cystic fibrosis; diagnosis; therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV55098
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:

1.55e-81 1017.00 96.3% 90.7% 91.1%

Length:
Matches:
Conservative:
Mismatches:

714 194 12 8 0

US-10-635-908-15 (1-214) x AAV55098

(1-714)

Indels: Gaps: Pred. 1 Score:

No . .

Alignment

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GlyGlnSerProLysLeuLeuIleTyrSerAlaSerAsnArgTyrThrGlyValProAsp

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This DNA sequence encodes a polypeptide (see AAW69307) comprising the CC light chain variable region of murine anti-interleukin-8 (II-8) CC monoclonal antibody (MAb) 5.12.14 (see AAW69305) and the human IgG1 light CC chain constant region. It was obtained by PCR amplification (see AAV55094 CC -95) and has been utilised in the construction of 5.12.14 Fab vector CC pantiII-8.2 (ATCC 97056). Humanised anti-II-8 MAbs are described for use in diagnostic applications and in the treatment of inflammatory CC disorders. The invention provides conjugates of an antibody fragment and a polymer, such as PEG, that have improved half-life, mean residence therapy of e.g. psoriasis, responses associated with inflammatory bowel disease (such as Crohn's disease associated with inflammatory bowel disease (such as Crohn's disease sundone, dermatitis, meningitis, concephalitis, uveitis, autoimmune diseases syndrome, dermatitis, meningitis, central nervous system inflammatory disorder, multiple organ injury CC syndrome secondary to septicaemia or trauma, alcoholic hepatitis, inflammations of the lung, including pleurisy, alveolitis, vasculitis, considerial pneumonia, chronic bronchitis, bronchiectasis, and cystic fibrosis.

CC (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                           New conjugates of antibody fragments - having covalently attached non-proteinaceous polymer molecules, particularly polyethylene glycol, for improving the residence time in the circulation.
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22-JAN-1998;
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colitis; immunotherapy; bacterial pneumonia; variable region;
light chain; heavy chain; neutrophil chemotaxis inhibitor;
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/product= "5.12.14-IgG1 construct"
/note= "5.12.14 light chain variable region
IgG1 light chain constant region construct"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence encodes the murine 5.12.14 (anti-IL-8) monoclonal antibody (Mab) light chain variable region and the human IgG1 light chain constant region construct. This construct is used in a novel method for treating ulcerative colitis which involves the administration of an anti-IL-8 monoclonal antibody capable of binding to human interleukin-8 (IL-8) with a Kd of 10-8 to 10-11 M. This Mab also inhibits neutrophil chemotaxis in response to IL-8, inhibits IL-8-mediated elastase release by neutrophils and does not bind to C5a, beta -TG or platelet factor 4. The anti-IL-8 antibodies can also be used for the treatment of bacterial pneumonia
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P-PSDB; AAW40122.
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LeuSerSerProValThrLysSerPheAsnArgGlyGluCys
                                                                                  LeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCysGluValThrHisGlnGly
                                                                                                                                        GAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGACG
                                                                                                                                                              GluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThr
                                                                                                                                                                                                                  CCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAG
                                                                                                                                                                                                                                        ProArgGluAlaLysValGlnTrpLysValAspAsnAlaLeuGlnSerGlyAsnSerGln
                                                                                                                                                                                                                                                                                                TCTGATGAGCAGTTGAAATCTGGAACTGCTTCTGTTGTGTGCCTGCTGAATAACTTCTAT
                                                                                                                                                                                                                                                                                                              SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr
                                                                                                                                                                                                                                                                                                                                                                          GGGACCAAGCTGGAGCTTCGAAGAGCTGTGGCTGCACCATCTGTCTTCATCTTCCCGCCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                  GAAGACTTGGCAGACTATTTCTGTCAGCAATATAACATCTATCCTCTCACGTTCGGTCCT
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                                                              CTGAGCAAAGCAGACTACGAGAAACACAAAAGTCTACGCCTGCGAAGTCACCCATCAGGGC
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20-FEB-1998;
24-JUL-1998;
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                                                                                                                              The present invention describes a novel conjugate having one or more antibody fragments covalently attached to one or more nonproteinaceous polymer molecules, where the apparent size of the conjugate is at least about 500 kDa. Conjugates of antibody fragments which bind the human interleukin (IL) 8 with a nonproteinaceous polymer can be used for treating inflammatory disorders e.g. acute lung injury, ischaemic reperfusion disorder, and autoimmune diseases. They can also be used for treating e.g. inflammatory skin diseases including psoriasis and atopic dermatitis, systemic scleroderma and sclerosis, and astimatic diseases. The conjugates can also be used as reagents in an animal model system for in vivo study of the biological functions of the antigen recognised by the conjugate. The present sequence encodes the 5.12.14 light chain variable region and the human IgGl light chain constant region fusion in the antigen recognised by the conjugate.
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                                                                                                                                                                                                                                                                                                                                         New conjugates of nonproteinaceous polymers for treating inflammatory disorders.
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98WO-US003337.
98US-00121952.
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Homo sapiens.
Chimeric.
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06-JUN-2000
                                                                                                                                                                                                                                                                                          Interleukin-8; IL-8; monoclonal antibody; MAb; anti-IL-8; 6G4.2.5V11N35A; inflammatory disorder; adult respiratory distress syndrome; mouse; affinity purification; 5.12.14; ds.
                      21-FEB-1997;
22-JAN-1998;
                                                                                  20-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                   Chimeric anti-IL-8 antibody light chain DNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an anti-interleukin-8 (IL-8) monoclonal antibody (MAb). The anti-IL-8 MAb comprises a sequence containing the CDRs (complementarity determining regions) of the humanized anti-IL-8 (G4.2.5ViIN35A light chain; and amino acids 24-253 of the humanized anti-IL-8 (G4.2.5ViIN35A heavy chain. The anti-IL-8 Abs and fragments can be used in diagnosis, for affinity purification of IL-8 from recombinant cell culture or natural sources and for the treatment of inflammatory disorders e.g. adult respiratory distress syndrome. Nucleic acids encoding the anti-IL-8 MAb can be associated in a vector with another gene encoding another protein or protein fragment to produce a fusion protein which can make isolation and/or purification of the protein an easier process. The present sequence represents the nucleotide sequence of murine anti-IL-8 antibody 5.12.14 light chain variable region and the human igg1 light chain constant region. (Updated on 15-SEP-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New nucleic acid molecule encodes a polypeptide which is an anti-
interleukin-8 monoclonal antibody or antibody fragment useful for the
production of anti-interleukin-8 monoclonal antibodies or fragments.
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Best Local Similari
US-10-635-908-15 (1-214) x AAC65491
                                                                                                                                                                                                                                                                                                                                     The present invention provides a number of humanised monoclonal anti-IL-8 antibodies which can be used in the diagnosis and treatment of inflammatory disorders, including adult respiratory distress syndrome, septic shock, multiple organ failure, bacterial pneumonia and inflammatory bowel disease. The present sequence encodes one of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Humanized anti-interleukin 8 monoclonal antibody variant useful treating inflammatory disorders, such as adult respiratory distresyndrome, hypovolemic shock and ulcerative colitis.
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22-JAN-1998;
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                        22-JAN-1998;
20-FEB-1998;
24-JUL-1998;
24-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                      Antibody; monoclonal antibody; 5.12.14; 6G4.2.5; interleukin-8; mAbj antiinflammatory; respiratory; acute lung injury; polyethylene glycol; pEG; lung injury; adult respiratory distress syndrome; ARDS; asthma; inflammatory disease; inflammatory bowel disease; psoriasis; sclerosis; ischaemic reperfusion disorder; stroke; multiple sclerosis; meningitis; osteoarthritis; septic shock; autoimmune disease; rheumatoid arthritis; alcoholic hepatitis; cystic fibrosis; ds; mouse; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mouse anti-IL-8 mAb 5.12.14 VL/human IgG1 light chain
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Synthetic.
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98US-0094003P.
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Percent Similarity:
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Query Match:
DB:

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Length:
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GTCACCTGCAAGGCCAGTCAGAATGTGGGTACTAATGTAGCCTGGT

GATATCGTCATGACACAGTCTCAAAAATTCATGTCCACATCAGTAGGAGACAGGGTCAGC AspIleValMetThrGlnSerGlnArgPheMetSerThrThrValGlyAspArgValSer

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ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer

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GluAspLeuAlaAspPhePheCysGlnGlnTyrSerAsnTyrProTrpThrPheGlyGly

81

190 41

GlyGlnSerProLysLeuLeuIleTyrSerAlaSerAsnArgTyrThrGlyValProAsp

GGGCAATCTCCTAAAGCACTGATTTACTCGTCATCCTACCGGTACAGTGGAGTCCCTGAT

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Comprising administering to the mammal an effective amount of a conjugate cof a single antibody fragment covalently attached to 1 or 2 polyethylene cof glycol (PEG) molecules, where the antibody fragment is a F(ab')_2 comprising: (a) first chain that is either a light chain or a heavy chain (c) if the first chain or a light chain opposite the covalently attached to 1 or 2 polyethylene comprising: (a) first chain that is either a light chain or a heavy chain; (b) a second chain that is either a light chain or a heavy chain; (c) a covalently attached to a first cysteine residue in the first cysteine residue in the first or second heavy chain; where covalently attached to a first cysteine residue in the first or second cysteine residue in the first or second dysteine residue in the first or second dysteine residue in the first or second opposite the second cysteine residue in the first or second opposite chain, where the first or second opposite chain, where the first or second opposite the first or second opposite chain, where the second cysteine residue in the first or second opposite the chain, where chain, where the first or second opposite the the binds to human interleukin-8 (II-8), and where the apparent size of the conjugate is at least about 500 kD. The antigen binding sites may be derived from continuing monoclonal antibodies 5.12.14 or 664.2.5. The method is useful for treating lung injury, including adult respiratory discress syndrome (ARDS) in a mammal and inflammatory diseases (such as asthma, continuing and continuing and sectores syndrome continuing and con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating acute lung injury in mammal by administering to mammal a 500 conjugate comprising F(ab') 2 antibody fragment that binds to human interleukin-8, covalently attached to one or two polyethylene glycol
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Search completed: June 3, 2006, 04:12:54 Job time: 708.553 secs

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Command line parameters:

-MODEL=frame+ p2n.model -DEV=xlh
-Q=|abse/BasSwEB_spool/UR10635908/runat_02062006_104216_10273/app_query.fasta_1
-Q=|abse/BasSwEB_spool/UR10635908/runat_02062006_104216_10273/app_query.fasta_1
-DB=EST -QFMT=fastap -SUFFIX=rst -MINMATCH=0.1 -LOOPEXT=0
-UNITS=blts -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi -LIST=45
-DOCALIGN=200 -THR_SCORESPOT -TRM_MAX=100 -THR_MIN=0 -ALIGN=15 -MODE-LOCAL
-OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=2000000000 -HOST=abse05h
-USER=US10635908_@CGN_1_1_7986_@runat_02062006_104216_10273 -NCFU=6 -TCFU=3
-NO_MMAP -NEG_SCORES=0 -MAIT -DSPBHOCK=100 -LONGLOG -DEV_TIMEOUT=120
-NO_MMAP -NEG_SCORES=0 -MAIT -DSPBHOCK=100 -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
-YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELLEXT=7
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Listing first 45 summaries
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Ygapop 10.0 , Ygapext
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Copyright (c) 1993 - 2006 Biocceleration Ltd
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## ALIGNMENTS

		JOURNAL	AUTHORS	REFERENCE			ORGANISM	KEYWORDS SOURCE	VERSION	ACCESSION	DEFINITION	LOCUS	RESULT 1
DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLCM1702 row: a column: 02 High quality sequence stop: 805.	Email: Ggapbs-remail.nin.gov Tissue Procurement: Louis M. Staudt, M.D., Ph.D. CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)	Unpublished (1999) Contact: Robert Strausberg, Ph.D.	National Institutes of Health, Mammalian Gene Collection (MGC)	1 (bases 1 to 867)	Hominidae; Homo.	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia: Eutheria: Euarchontoglires; Primates; Catarrhini;	Homo sapiens	Homo sapiens (human)	BG754732.1 GI:14065385	mRNA sequence. BG754732	1F1 NIH_MGC_48 Hc	BG754732 867 bp mRNA linear EST 15-MAY-2001	

linear

EST 15-MAY-2001

Gene Collection (MGC)

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                               SerSerProValThrLysSerPheAsnArgGlyGluCys 214
                                                                                                                           SerLysAlaAspTyrGluLysHisLysValTyrAlaCysGluValThrHisGlnGlyLeu
                                                                                                                                                                                                                                 SerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThrLeu
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  AGCTCGCCCGTCACAAAGAGCTTCAACAGGGGAGAGTGT
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/clone="MAGE:4854409"
/tissue_type="primary B-cells from tonsils (cell line)"
/tissue_type="primary B-cells from tonsils (cell line)"
/lab host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_48"
/clone_lib="NIH_MGC_48"
/clone_Torgan: B-cells; Vector: poTB7; Site_1: XhoI;
/site_2: EcoRI, cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGA(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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/mol_type="mRNA"
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Tissue Procurement: Louis M. Staudt,
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National Institutes of Health, Mammalian
Unpublished (1999)
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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CDNA Library Preparation: Ling Hong/Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMstGlnSer
                                                                            GGGAAAGCCCCTAAGCTGCTAATCTACGATGCATCTAATTTGGAAACAGGGGTCCCATCA
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/lab_host="primary B-cells from tonsils (cell line)"
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/db_xref="taxon:9606"
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Clone distribution: MGC clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLCM1705 row: 1 column: 16 High quality sequence stop: 740.
Location/Onalifian-
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BG756519.1 GI:
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602715662F1 NIH_MGC_48
                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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National Institutes of Health, Mammalian
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Mammalia; Eutheria;
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/tissue_type="primary B-cells from tonsils (cell line)"
/tlab host="DH10B (phage-resistant)"
/clone_lib="NIH MGC_48"
/note="Togan: B-cells; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(6). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of
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/db_xref="taxon:9606"
                                                                                                                               'clone="IMAGE:4855839"
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California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologion Note: this is a NIH\_MGC Library." Technologies).

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                   Homo sapiens
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Mammalia; Eutheria;
           Hominidae; Homo.
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Tissue Procurement: Dr. Mark Watson
CDNA Library Preparation: Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution: MGC clone distribution: MGC clone distribution for through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLCM2385 row: p column: 05
High quality sequence stop: 697.
Location/Qualifiers
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National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                           CCATCTGATGAGCAGTTGAAATCTGGGAACTGCCTCTGTTGTGTGCCTGCAATAACTTC
                                                                                                                                                                           GlyGlyThrLysLeuGluIleLysArgThrValAlaAlaProSerValPheIlePhePro
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  TATCCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCC
                         TyrProArgGluAlaLysValGlnTrpLysValAspAsnAlaLeuGlnSerGlyAsnSer
                                                                                                  ProSerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPhe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:6216052"
/lab_host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_113"
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/mol_type="mRNA"
/db_xref="taxon:9606"
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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/db_xref="taxon:906"
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/clone="IMAGE:3035409"
/clone=lib="NHIB (T1 phage-resistant)"
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US-10-635-908-15 (1-214) x CB956774 (1-810)

Query Match:

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         Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLCM1208 row: j column: 02
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1 (bases 1 to 943)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian

Unpublished (1999)
                                                                                                                                                                                                                                                                                                        mRNA sequence.
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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quality sequence stop: 721.
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s cDNA clone IMAGE:4336225 5',
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    LeuSerSerProValThrLysSerPheAsnArgGlyGluCys
                                                                                                                            GAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGACG
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                                                                                 LeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCysGluValThrHisGlnGly
                                                                                                                                            CCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCCCTCCAATCGGGTAACTCCCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone="IVMAGE:4336225"
/tissue_type="primary B-cells from tonsils (cell line)"
/lab host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_48"
/clone_lib="NIH_MGC_48"
/clone_lib="NIH_MGC_48"
/note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
/site_2: EooRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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5', mRNA sequence.
BQ706786
BQ706786.1 GI:21845685
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://imagge.lnl.gov
Plate: LLCM2382 row: o column: 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. Mark Watson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Euarchontoglires; Primates; Catarrhini;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1999)
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                     ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer 80
                                                                                                                 GlyGlnSerProLysLeuLeuIleTyrSerAlaSerAsnArgTyrThrGlyValProAsp
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Location/Qualifiers
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/lab_host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_113"
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/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                            2 rue Gaston Cremieux, CP 5706 - 91057 EVRY cedex - FRANCE Email: segref@genoscope.cns.fr, Web: www.genoscope.cns.fr
Email: segref@genoscope.cns.fr, Web: www.genoscope.cns.fr
1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-strand cDNA was digested with Not I and cloned into the Not I and EcoR V sites of the pcMVSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies, a division of Invitrogen. This sequence belongs to sequence cluster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hominidae; Homo.

1 (bases 1 to 774)

Li,W.B., Gruber,C., Jessee,J. and Polayes,D.

Pull-length cDNA libraries and normalization

Unpublished (2001)
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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774 bp mRNA linear EST 08-APR-200
bX336281 Homo sapiens PLACENTA COT 25-NORMALIZED Homo sapiens cDNA
clone CS0DI026YL22 5-PRIME, mRNA sequence.
                                                                                                                                                                                                                                                                                                  For more information about this cluster, see http://www.genoscope.cns.fr/cdna?s=CSODIO26DF11QP1&c=1696.r. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               On May 2, 2003 this sequence version replaced gi:30341499. Contact: Genoscope
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/tissue type="PIACENTA COT 25-NORMALIZED"
/clone_Tib="Homo sapiens PLACENTA COT 25-NORMALIZED"
/note="Ist strand cDNA was primed with a NotI-oligo(dT)
primer. Five prime end enriched, double-strand cDNA was
digested with Not I and cloned into the Not I and Ecor v
sites of the pCMVSPORT 6 vector. Library was normalized.
                                                                                                                                                                  /db_xref="taxon:9606"
/clone="CSODI026YL22"
                                                                                                                                                                                                                                               organism="Homo sapiens"
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5', mRNA sequence.
BQ707105
BQ707105.1 GI:21846004
EST.
Hominidae; Homo.

1 (bases 1 to 925)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Unpublished (1999)

Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Dr. Mark Watson
                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                     Homo sapiens (human)
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cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Agencourt Bioscience Corporation

Clone distribution: MGC clone distribution information can

found through the I.M.A.G.E. Consortium/LLNL at:
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plate: LLCM2473 row: g column: 10
High quality sequence stop: 647.
Location/Qualifiers
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                               GluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThr 180
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/clone_lib="NHH_MGC_113"
/clone_lib="NHH_MGC_113"
/note="Organ: spleen; Vector: pOTB7; Site_1: XhoI; Site_2 note="Organ: spleen; Vector: pOTB7; Site_1: XhoI; Site_2 note="Organ: spleen; Vector: pOTB7; Site_1: XhoI; Site_2 note="Organ: spleen; Sites note="
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:6281121"
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LeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCysGluValThrHisGlnGly

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1 (bases 1 to 803)

1 (bases 1 to 803)

NIH-MGC thttp://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Dr. Michael Brownstein and Dr. Miklos Palk

cDNA Library Preparation: CLONNTSCH Laboratories, Inc.

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Agencourt Bioscience Corporation

Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CB958596
CB958596.1 GI:30214712
EST.
Homo sapiens (human)
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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                                          e: NDCM155 row: o column: 11 quality sequence stop: 667. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                              /clone="IMAGE:30354106"
//lab host="DH10B (T1 phage-resistant)"
//lab host="DH10B (T1 phage-resistant)"
//clone_lib="NIH_MGC_184"
//clone_lib="NIH_MGC_184"
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//clone="NIH_MGC_184"
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/mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
CDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLCM1707 row: a column: 24
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602716389F1 NIH_MGC_48 Homo sapiens cDNA clone IMAGE:4856351 5',
mRNA seconence
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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                                                                                                                                   quality sequence stop:
/organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:

1.31e-96 924.00 89.7% 81.3% 82.8%

Score: Pred. No.: Alignment ORIGIN

Scores:

US-10-635-908-15 (1-214) x BG755921

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BG755394 750 bp r 602713951F1 NIH_MGC_48 Homo sapiens mRNA sequence.
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/lab_host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_48"
/clone_lib="NIH_MGC_48"
/note="Porgan: B-cells; Vector: pOTB7; Site_1: XhoI;
Site_2: ECORI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the
following 5' adaptor: GGCACAGG(G). Size-selected >500bp
for average insert size 1.8kb. Library constructed by Ling
Hong in the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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Tissue Procurement: Louis M. Staudt, M.D., Ph.D.
CDNA Library Preparation: Ling Hong/Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1701 row: e column: 14
High quality sequence stop: 711.
location/Qualifiers
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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Mammalia; Eutheria;
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/db xref="taxon:9606"
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/clone="IMAGE,4854133"
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/tissue_type="primary B-cells from tonsils from tonsils from tonsils from tonsils from tonsils from tonsils from the strength from the laboratory of GGCACGAG(G). Size-selected by Ling for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of Galfornia, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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Contact: Robert Strausberg, Ph.D.
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National Institutes of Health, Mammalian
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Mammalia; Eutheria;
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                                              /clone="IMAGE:4854374"
/tlssue_type="primary B-cells from tonsils (cell line)"
/tlssue_type="primary B-cells from tonsils (cell line)"
/lab host="DH10B (phage-resistant)"
/clone_1ib="NIH MGC 48"
/clone_1ib="NIH MGC 48"
/note="Organ: B-cells; Vector: pOTB7; Site_1: XhoI;
Site_2: EccRI; cDNA made by oligo-dT priming.
Site_2: EccRI; cDNA made by ling for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
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National Institutes of Health, M
Unpublished (1999)
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AGENCOURT_6940425 NIH_MGC_99
5', mRNA sequence.
BQ056652
 Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov Tissue Procurement: Lou Staudt
                                                                                                                                          Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                   Homo sapiens
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cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
cNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can l
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM2054 row: c column: 02
High quality sequence start: 13
High quality sequence stop: 695.
Location/Qualifiers
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GluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThr 180
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/clone="IMAGE:5809081"
/tlssue type="lymphoma, cell line"
/tlssue type="NIH_MGC_9"
/clone_lib="NIH_MGC_9"
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cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
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High quality sequence start: 23
High quality sequence stop: 711.
Location/Qualifiers
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Tissue Procurement: Lou Staudt
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1 (bases 1 to 1047)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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EST.
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BQ059853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LeuSerSerProValThrLysSerPheAsnArgGlyGluCys 214
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AspIleValMetThrGlnSerGlnArgPheMetSerThrThrValGlyAspArgValSer 20
                                                                                                                                                                                                                                                                                                                                          /tissue_type="lymphoma, cell line"
/tissue_type="lymphoma, cell line"
/lab host="DH10B (phage-resistant)"
/clone lib="NIH MGC 99"
/clone lib="NIH MGC 99"
/note="Organ: lymph; Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Organ: lymph; Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Corgan: lymph; Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Corgan: lymph; Vector: pOTB7; Site_1: XhoI; Cloned
into EcoRI/XhoI sites using the following 5: adaptor:
GGCACGAG(G). Size-selected >500bp for average insert size
1.8kb. Library constructed by Ling Hong in the laboratory
of Gerald M. Rubin (University of California, Berkeley)
using ZAP-cDNA synthesis kit (Stratagene) and Superscript
IRT (Life Technologies). Note: this is a NIH_MGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mol_type="mRNA"
/db_xref="taxon:9606"
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Matches:
Conservative:
Mismatches:
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                                                                                                                                                                                    SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr 140
                                                                                                                                                                                                                  GlyThrLysLeuGluIleLysArgThrValAlaAlaProSerValPheIlsPheProPro 120
                                                                                                                                                                                                                                                                                                        ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer 80
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AGGTTCAGCGGCAGTGGATATGGGACAGAATTCAGTCTCACCATCAGCAGCCTGCAGCCT 354
                                         ProArgGluAlaLysValGInTrpLysValAspAsnAlaLeuGlnSerGlyAsnSerGln 160
            LeuSerSerProValThrLysSerPheAsnArgGlyGluCys
                                                                                                                                                                         TCTGATGAGCAGTTGAAATCTGGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTAT
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                    214
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Search completed: June 3, 2006, 09:16:28 Job time: 5400.06 secs

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Command line parameters:

-MODEL-frame+ p2n.model -DEV=xlh
-Q=/abss/ABSSWEB spool/US10635908/runat 02062006 104220 10327/app query.fasta_1
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-DB=Issued Patents NA -QrMT=fastap -SUFFIX=rni -MINMATCH=0.1 -LOOPCL=0
-DB=ISSUED Patents NA -QrMT=fastap -SUFFIX=rni -MINMATCH=0.1 -TRANS=human40.cdi
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=bitsusman 2-TRANS=human40.cdi
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-DEV TIMEOUT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                          Score
                                 1017
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1: /EMC_Celerra_SIDS3/ptodata/2/ina/5_COMB.seq:*

2: /EMC_Celerra_SIDS3/ptodata/2/ina/6A_COMB.seq:*

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5: /EMC_Celerra_SIDS3/ptodata/2/ina/7_COMB.seq:*

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10: /EMC_Celerra_SIDS3/ptodata/2/ina/Backfiles1.seq:*
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Ygapop 10.0 , 1
Fgapop 6.0 , 1
Delop 6.0 , 1
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                                                                                                                                                          Match Length
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                                                                                                                                                                            Query
 DIVMTQSQRFMSTTVGDRVS......EVTHQGLSSPVTKSFNRGEC 214
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Ygapext
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Delext
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US-08-398-613A-27

US-08-398-611A-27

US-08-398-611A-27

US-08-396-851A-27

US-08-491-334A-27

US-08-804-444A-24

US-08-804-444A-24

US-09-026-985-24
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 Sequence 27,
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Sequence 27,
Sequence 27,
Sequence 27,
Sequence 24,
Sequence 24,
Sequence 24,
                                                                                                                                                            Description
 US-08-398-613A-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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## ALIGNMENTS

Hebert, Caroline Alice

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COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 Kb flopp
MEDIUM TYPE: 5.25 inch, 360 Kb flopp
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/398,613A
FILING DATE: 01-MAR-1994
                  CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/2
FILING DATE: 03-MAR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Fong, Sherman
APPLICANT: Hebert, Carolli
APPLICANT: Kim, Kyung Jin
APPLICANT: Leong, Steven I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Leong, STITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                   STREET: 460 Point San Brun
CITY: South San Francisco
STATE: California
COUNTRY: USA
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                                                                                                                                                                                                                                                                                                         94080
Fitte,
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                                                                08/205864
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REGISTRATION NUMBER: 35,136
REFERENCE/DOCKET NUMBER: 874F1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/25-1489
TELEFAX: 415/25-1489
TELEFAX: 415/25-1981
TELEY 910/371-7168
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERIST
RESULT 2
US-08-398-612A-27
i Sequence 27, Application US/08398612A
; Patent No. 5686070
; GENERAL INFORMATION:
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Percent Similarity:
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Query Match:
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APPLICATION NUMBER: 08/398611
FILING DATE: 01-Mar-1995
APPLICATION UNMBER: 08/205864
FILING DATE: 03-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: LOVE, RICHARD B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: P0874P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5530
TELEPHONE: 415/225-9881
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb flopp)
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/398,612A
FILING DATE: 01-MAR-1995
CLASSIFICATION: 424
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APPLICANT: Fong, Sherman
APPLICANT: Hebert, Caroline Alice
APPLICANT: Kim, Kyung Jin
APPLICANT: Leong, Steven R.
TITLE OF INVENTION: Anti-IL-8 Monoclonal Antibodies for
TITLE OF INVENTION: Treatment of Inflammatory Disorders
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TYPE: Nucleic Acid
STRANDEDNESS: Double
TOPOLOGY: Linear
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CITY: South San Francisco
STATE: California
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US-08-398-611A-27
                                   APPLICATION NUMBER: 08/205864
PILING DATE: 03-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: LOVE, RICHARD 8
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: P0874P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-5530
TELEPHONE: 415/225-9881
TELEPAX: 415/952-9881
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 714 base pairs
TYPE: Nucleic Acid
CTE NUCLEIC ACID
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MEDIUM TYPE: 3.5 inch, 1.44 Mb flopp:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/398,611A
FILING DATE: 01-Mar-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
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APPLICANT: Doersh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Leong, Steven R.
TITLE OF INVENTION: Anti-IL-8 Monoclonal Antibodies for
TITLE OF INVENTION: of Inflammatory Disorders
NUMBER OF SEQUENCES: 58
CORRESPONDENCE ADDRESS:
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APPLICANT: Hebert, Caroline Alice
APPLICANT: Kim, Kyung Jin
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STREET: 460 Point San Bruno
CITY: South San Francisco
STATE: California
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                          STRANDEDNESS:
               TOPOLOGY:
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                                                                                    GENERAL INFORMATION:
APPLICANT: Fong, Sherman
APPLICANT: Hebert, Caroline Alice
APPLICANT: Kim, Kyung Jin
APPLICANT: Kim, Kyung Jin
APPLICANT: Leong, Steven R.
TITLE OF INVENTION: Anti-IL-8 Monoclonal Antibodies
NUMBER OF SEQUENCES: 58
CORRESPONDENCE ADDRESS:
   STREET: 460 Point San Bruno Blvd CITY: South San Francisco STATE: California COUNTRY: USA
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Percent Similarity:
Best Local Similarity:
Query Match:
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Pred. No.:
US-08-396-851A-27; Sequence 27, Application; Patent No. 5707622
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Query Match:
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MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UMMBER: US/08/396,851A
FILING DATE: 01-MAR-1995
CLASSIFICATION 1424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/205864
FILING DATE: 03-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Fitte, Renee A.
REGISTRATION NUMBER: 35,136
REGISTRATION INFORMATION:
TELEPHONE: 415/225-1489
TELEPHONE: 415/225-1489
TELEPHONE: 415/225-1489
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 714 bases
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STRANDEDNESS: single
TOPOLOGY: linear
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GluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThr 180
                                                                           ProArgGluAlaLysValGlnTrpLysValAspAsnAlaLeuGlnSerGlyAsnSerGln 160
                                                                                                                                       SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr 140
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                                                                                                                       TCTGATGAGCAGTTGAAATCTGGAACTGCTTCTGTTGTGTGCCTGCTGAATAACTTCTAT
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                                                                                                                                                                                                                     Percent Similarity:
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COMPUTER READABLE FORM:

MEDIUM TYPE 3.5 inch, 1.44 Mb floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/491,334A

FILING DATE: 27-Jun-1995

CLASSIFICATION DATA:

APPLICATION UMBER: 08/398611

FILING DATE: 01-MAR-1995

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/20864

FILING DATE: 03-MAR-1994

ATTORNEY/AGENT INFORMATION:

NAME: LOVE, RICHARD B.

REGISTRATION NUMBER: 34,659

REGISTRATION INFORMATION:

TELEPHONE: 415/225-550

TELEPHONE: 415/225-550

TELEPHONE: 415/225-9881
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Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
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APPLICANT: Kabakoff, Rhona C.
APPLICANT: Moore, Mark W.
TITLE OF INVENTION: IL-8 Antagonists for Treatment of Inflammatory
TITLE OF INVENTION: Disorders and Asthma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 460 Point San Bruno
CITY: South San Francisco
STATE: California
COUNTRY: USA
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STRANDEDNESS: Double
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                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 714 base pairs
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21 IleThrCysLysAlaSerGlnAsnValValSerAlaValAlaTrpTyrGlnGlnLysPro 40
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Matches:
Conservative:
Mismatches:
Indels:
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RESULT 6
US-09-027-449-24
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Patent No. 6
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MEDIUM TYPE: 3.5 inch, 1.44 mb floppy di
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/027,449
FILING DATE: 20-Feb-1998
                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Gonzalez, Tania R.
APPLICANT: Gonzalez, Tania R.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Antibody Fragment-Polymer Conjugates
TITLE OF INVENTION: Humanized Anti-IL-8 Monoclonal Antibo
NUMBER OF SEQUENCES: 72
CORRESPONDENCE ADDRESS:
FILING DATE: 20-Feb-1998
CCLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/074,330
FILING DATE: 22-Jan-1998
PRIOR APPLICATION NUMBER: 60/038,664
APPLICATION NUMBER: 60/038,664
FILING DATE: 21-Feb-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                            STREET: 1 DNA WAY
CITY: South San Francisco
STATE: California
COUNTRY: USA
ZIP: 94080
ZIP: 94080
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   RESULT 7
US-08-04-444A-24
; Sequence 24, Application US/08804444A
; Patent No. 6117980
; GENERAL INFORMATION:
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Query Match:
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REGISTRATION NUMBER: 91085R3-2
REFERENCE/DOCKET NUMBER: P1085R3-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/925-5530
TELEPHONE: 650/925-9881
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 714 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Double
STRANDEDNESS: Double
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TELEPHONE: 650/225-530
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 714 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Double
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy di
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/804,444A
FILLNG DATE: 21-Feb-1997
CLASSIFICATION: 530
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Leong, Steven R.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Humanized Anti-IL-8 Monoclonal Antibodies
NUMBER OF SEQUENCES: 61
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: LOVE, RICHARD B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: P1085
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CITY: South San Francisco
STATE: California
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                  SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr 140
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy dis

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTMARE: WinPatin (Genentech)

CURRENT APPLICATION NUMBER: US/09/026,985

FILING DATE: 20-Peb-1998

CLASSIFICATION:

APPLICATION NUMBER: 34,659

REFERENCE/DOCKET NUMBER: 34,659

REFERENCE/DOCKET NUMBER: 91085R3-1

TELECOMMUNICATION INFORMATION:

TELEPAX: 650/225-530

TELEPAX: 650/952-9881

INFORMATION FOR SEQ ID NO: 24:

SEQUENCE CHARACTERISTICS:

LENGTH: 74 base pairs
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US-09-026-985-24
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                                                                                                                  Percent Similarity:
Best Local Similarity:
Query Match:
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US-09-026-985-24
                                                                 US-10-635-908-15 (1-214)
                                                                                                                                                                       Score:
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GENERAL INFORMATION:
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APPLICANT: Leong, Steven R.
APPLICANT: Leong, Steven R.
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: Antibody Fragment-Polymer Conjugates and
TITLE OF INVENTION: Humanized Anti-IL-8 Monoclonal Antibodies
NUMBER OF SEQUENCES: 72
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                         No.:
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STATE: California
                                                                                                                                                                                                                                                                       LENGTH: 714 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Double
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Application US/09026985
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Mismatches:
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COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/121,952A

FILING DATE: 24-Jul-1998

CLASSIFICATION 1514

PRIOR APPLICATION DATA:

APPLICATION UMBER: 60/074330

FILING DATE: 22-JAN-1998
                                                                                                                                                                                                                                                                             APPLICANT: Genentech, Inc., Hsei, Vanessa
APPLICANT: Koumenis, Iphigenia
APPLICANT: Koumenis, Iphigenia
APPLICANT: Leong, Steven R.
APPLICANT: Leong, Steven R.
APPLICANT: Presta, Leonard G.
APPLICANT: Shahrokh, Zahra
APPLICANT: Shahrokh, Zahra
APPLICANT: Zapata, Gerardo A.
APPLICANT: Zapata, Gerardo A.
TITLE OF INVENTION: WITH ANTI-IL-8 ANTIBODY FRAGMENT-POLYMER CONJUGATES
NUMBER OF SEQUENCES: 72
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APPLICATION NUMBER: 60/075467
FILING DATE: 20-FEB-1998
ATTORNEY/AGENT INFORMATION:
NAME: LOVE, RICHARD B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: P108:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/25-530
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 714 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Double
TOPDICGY: Tipear
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                                                        LeuSerSerProValThrLysSerPheAsnArgGlyGluCys
                                       CTGAGCTCGCCCGTCACAAGAGCTTCAACAGGGGAGAGTGT
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RESULT 9 US-09-121-952A-24

Patent No. 6458355

Sequence 24, Application US/09121952A

ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
COUNTRY: USA
ZIP: 94080

CORRESPONDENCE ADDRESS

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670 201 610

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Query
DB:
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                 Pred. No.:
                                                                                                                                                                                                                                                                                                                                  Alignment Scores:
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 94080
COMPUTER READABLE PORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/234,340A
FILING DATE:
CLASSIFICATION: COMPANIEST CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/121,95

FILING DATE: 24-Jul-1998

APPLICATION NUMBER: 60/074330

FILING DATE: 22-JAN-1998

PRIOR APPLICATION DATA:
APPLICATION DATA: 60/075467

FILING DATE: 20-FEB-1998

ATTORNEY/AGENT INFORMATION:
NAME: LOVE, RICHARD B.

REFERENCE/DOCKET NUMBER: 91,659

REFERENCE/DOCKET NUMBER: D1085R4

TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 650/952-9881 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Genentech, Inc., Hsei, Vanessa
APPLICANT: Koumenis, Iphigenia
APPLICANT: Koumenis, Iphigenia
APPLICANT: Leong, Steven R.
APPLICANT: Presta, Leonard G.
APPLICANT: Shahrokh, Zahra
APPLICANT: Shahrokh, Zahra
APPLICANT: Zapata, Gerardo A.
APPLICANT: Zapata, Gerardo A.
APPLICANT: Zapata, Gerardo A.
APPLICANT: Zapata, Gerardo A.
APPLICANT: MINUNION: MITHOUS OF TREATING INFLAMMATORY DISEASES
TITLE OF INVENTION: WITH ANTI-IL-8 ANTIBODY FRAGMENT-POLYMER CONJUGATES
NUMBER OF SEQUENCES: 72
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LENGTH: 714 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Double
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STREET: 1 DNA Way
CITY: South San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
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                                                              130 GTCACCTGCAAGGCCAGTCAGAATGTGGGTACTAATGTAGCCTGGTATCAACAGAAACCA
190
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                                                                                          21 IleThrCysLysAlaSerGlnAsnValValSerAlaValAlaTrpTyrGlnGlnLysPro 40
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                                                                                                                              4, Application US/09234340A
6468532
                          GlyGlnSerProLysLeuLeuIleTyrSerAlaSerAsnArgTyrThrGlyValProAsp 60
GGCAATCTCCTAAAGCACTGATTTACTCGTCATCCTACCGGTACAGTGGAGTCCCTGAT
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91.1%
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Matches:
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US-09-355-014-24
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Sequence 24, Application US/09355014 Patent No. 6870033 GENERAL INFORMATION:
TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION NUMBER: US/09/355,014
PILING DATE: 21-Jul-199
CLASSIFICATION : <Unknown>
APPLICATION : <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Love, Richard B.
REGISTRATION NUMBER: 34,659
REFERENCE/DOCKET NUMBER: 91085R3PCT
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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Koumenis, Iphigenia
Leong, Steven R.
Presta, Leonard G.
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Zapata, Gerardo A.
TITLE OF INVENTION: Antibody Fragment-Polymer Conjugates and
Humanized Anti-IL-8 Monoclonal Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 72
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Genentech,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Inc., Hsei,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vanessa
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TYPE: Nucleic Acid STRANDEDNESS: Double TOPOLOGY: Linear

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Best

Match:

130

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Percent Similarity: Best Local Similarity:

1.86e-112 1017.00 96.3% 90.7% 91.1%

Alignment Scores:

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RESULT 12
US-09-726-258-24
(Sequence 24, Application US/09726258
; Sequence 24, Application US/09726258
; Patent NO. 7005504
; Patent NO. 7005504
; APPLICANT: Genentech, Inc., Hsei, Vanessa
; APPLICANT: Koumenis, Iphigenia
; APPLICANT: Leong, Steven R.
; APPLICANT: Leong, Steven R.
; APPLICANT: Shahrokh, Zahra
; TITLE OF INVENTION: ANTIBODY FRAGMENT-POLYMER CONJUGATES AND
; TITLE OF INVENTION: HUMANIZED ANTI-IL-8 MONOCLONAL ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-10-635-908-15 (1-214) x US-09-355-014-24 (1-714)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TCTGATGAGCAGTTGAAATCTGGAACTGCTTCTGTTGTGTGCCTGCTGAATAACTTCTAT
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PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/094003

RILING DATE: 24-JUL-1998

ATTORNEY/AGENT INFORMATION:

NAME: LOVE, RICHARDE B.

REGISTRATION NUMBER: 34.659

REFERENCE/DOCKET NUMBER: P108

TELEPHONE: 650/225-5530

TELEPAN: 650/952-9881

INFORMATION FOR SEQ ID NO: 24:

SEQUENCE CHARACTERISTICS:

LENGTH: 714 base pairs

TYPE: Nucleic Acid

STRANDEDNESS: Double

TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb fl.
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Windatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/726,258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 72
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, I
STREET: 1 DNA Way
CITY: South San Francis
STATE: Călifornia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                                        370
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                                                                                                                                                                                                                                                         GlyGlnSerProLysLeuLeuIleTyrSerAlaSerAsnArgTyrThrGlyValProAsp
                                                                                                                        GluAspLeuAlaAspPhePheCysGlnGlnTyrSerAsnTyrProTrpThrPheGlyGly 100
     SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr 140
                                                           ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer
                                         GGGACCAAGCTGGAGGCTTCGAAGAGCTGTGGCTGCACCATCTGTCTTCATCTTCCCGCCA
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90.7%
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Matches:
Conservative:
Mismatches:
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APPLICANT: Chapman, Andrew Paul
APPLICANT: King, David John
TITLE OF INVENTION: Biological Products
FILE REFERENCE: Carp-0089
CURRENT APPLICATION NUMBER: US/09/875,221B
CURRENT FILING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: GB0013810.7
PRIOR FILING DATE: 2000-06-06
NUMBER OF SEQ ID NOS: 130
SOFTWARE: Patentin version 3.1
SEQ ID NO 127
LENGTH: 642
                                                                                                                                                                                                                                                                                                                                                                                                Percent Similarity:
Best Local Similarity:
Query Match:
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US-09-875-221B-127
                                                                                                                                                                                                                                                                                                                                             US-10-635-908-15 (1-214) x US-09-875-221B-127 (1-642)
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Athwal, Dilject Singh
APPLICANT: Brown, Derek Thomas
APPLICANT: Weir, Andrew Neil Charles
APPLICANT: Popplewell, Andrew George
APPLICANT: Chapman, Andrew Paul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                      GlyGlnSerProLysLeuLeuIleTyrSerAlaSerAsnArgTyrThrGlyValProAsp
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GluAspLeuAlaAspPhePheCysGlnGlnTyrSerAsnTyrProTrpThrPheGlyGly 100
                                                           ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer
                                                                                                                 GGTAAAGCCCCAAAAAGCCCTCATCTACAGTGCCTCTTTCCTCTATAGTGGTGTACCATAC
                                                                                                                                                                                            ATCACTTGTAAAGCCAGTCAGAACGTAGGTACTAACGTAGCCTGGTATCAGCAAAAACCA
                                     AGGITCAGCGGATCCGGTAGTGGTACTGATTTCACCCTCACGATCAGTAGCCTCCAGCCA 240
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US-08-030-175-3
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Patent No. 6767996
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Gorman, Scott D.
APPLICANT: Clark, Michael R.
APPLICANT: Cobbold, Stephen P.
APPLICANT: Waldmann, Herman
TITLE OF INVENTION: ALTERED ANTIBODIES AND THEIR PREPARATION
NUMBER OF SEQUENCES: 43
                                                                                                                                                                                    APPLICATION UMBER: US/08/030,175
FILING DATE: 17-MAY-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: PCT/GB91/01578
FILING DATE: 13-SEP-1991
ATTORNEY/AGENT INFORMATION:
NAME: Ernst, Barbara G.
REGELSTRATION NUMBER: 30,377
REFERENCE/DOCKET NUMBER: 1768-113
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)/83-6040
                                                                                                                                  TELEFAX: (202)783-6031
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
               TOPOLOGY: linear MOLECULE TYPE: CDN
                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk, 5.25 inch, 3
COMPUTER: IT COMPATIBLE
COMPUTER: TO COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS V 3.2
SOFTWARE: Wordberfect 5.0 (Dos Text)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: D. C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Rothwell, Figg, Ernst & F
STREET: 555 13TH ST., NW Suite 701
                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U.S. ZIP: 20004
                                                                                                                   LENGTH:
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                                                                                             nucleic acid
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Best Local Similarity:
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US-08-952-235-3
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                                                                                                                               Sequence 3, Application US/08952235
Patent No. 6207152
                      GENERAL INFORMATION:
APPLICANT: Schwall, Ralph H.
APPLICANT: Tabor, Kelly H.
APPLICANT: Tabor, Kelly H.
TITLE OF INVENTION: Hepatocyte Growth Factor Receptor
TITLE OF INVENTION: Antagonists and Uses Thereof
NUMBER OF SEQUENCES:
ADDRESSE: Genetech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            No.:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer 80
                                                                                                                                                                                                                                                                                                                                                                                                                SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr 140
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                                                                                                                                                                                                              LeuSerSerProValThrLysSerPheAsnArgGlyGluCys 214
                                                                                                                                                                                                                                                                                                                        GluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThr
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                                                                                                                                                                                                  CTGAGCTCGCCGTCACAAAGAGCTTCAACAGGGGAGAGTGT
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            E: Genentech, Inc.
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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Best Local Similarity:
Query Match:
DB:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/952,
FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION A24
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/460368
FILING DATE: 02-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: MATSCHAPG, DIANE L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: 90931
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/25-9416
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 650 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: Califo
COUNTRY: USA
ZIP: 94080
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                                                                                                                                                                                                         241 ATCACCAGTGTGAAGGCTGACGACCTGGCAGTTTATTACTGTCAGCAATATTATGCCTAT
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GlnSerGlyAsnSerGlnGluSerValThrGluGlnAspSerLysAspSerThrTyrSer 174
                             ValPheIlePheProProSerAspGluGlnLeuLysSerGlyThrAlaSerValValCys
                                                                                                                                                               ProTrpThrPheGlyGlyGlyThrLysLeuGluIleLysArgThrValAlaAlaProSer 114
                                                                                         GTCTTCATCTTCCCGCCATCTGATGAGCAGTTGAAATCTGGAACTGCCTCTGTTGTGTGC
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943.00
92.3%
80.9%
84.5%
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154

420 134 360 300

240 74 180 54 120

175 LeuSerSerThrLeuThrLeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCys 194	Search completed: June 3, 2006, 06:53:38 Job time : 219.357 secs
175 LeuSerSerThrLeuThrLeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCys 194	CACCCATCAGGGCCTGAGCTCG
175 LeuserserThrleuThrleuserLysAlaAspTyrGluLysHisLysValTyrAlaCys 194	lThrHisGlnGlyLeuSerSer
175 LeuSerSerThrLeuThrLeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCys 194	CAGCACCCTGACGCTGAGCAAA
	rSerThrLeuThrLeuSerLys
481 CARTCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGGACAGGACAGCAACGTACAGC 540	GGGTAACTCCCAGGAGAGTGTC

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Run on:
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-MODEL-frame+ p2n.model -DEV=xlh
-MODEL-frame+ p2n.model -DEV=xlh
-Q=/abse/ABSSWEB_spool/US10635908/runat_02062006_104222_10380/app_query.fasta_1
-Q=/abse/ABSSWEB_spool/US10635908/runat_02062006_104222_10380/app_query.fasta_1
-DB=Published_App_lications_NA_Main -QFMT=fastap_-SUFFIX=rnpbm -MINMATCH=0.1
-LOOPECL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -AATRIX=blosum62
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -AATRIX=blosum62
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR_SCORE-PC-T-THR_MAX=100
-THR_MIN=0 -ALIGN=15 -MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXLEN=200000000 -HOST=abss02h
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -USER=US10635908 @CGN 1 1 2326 @runat 02062006 104222 10380 -NCPU=6 -ICPU=3 -NO_MMAP -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG -DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
                                                                                                                                                                                                                                                 pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                        Score
1017
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1: /EMC_Gelerra_SIDS3/ptodata/2/pubpna/US08_PUBCOMB.seq:*

2: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US08_PUBCOMB.seq:*

3: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US09A_PUBCOMB.seq:*

4: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US09B_PUBCOMB.seq:*

5: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US09C_PUBCOMB.seq:*

6: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

7: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US10B_PUBCOMB.seq:*

8: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US10C_PUBCOMB.seq:*

9: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US10E_PUBCOMB.seq:*

10: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US10E_PUBCOMB.seq:*

11: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US10E_PUBCOMB.seq:*

12: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US10E_PUBCOMB.seq:*

13: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US11A_PUBCOMB.seq:*

14: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US11A_PUBCOMB.seq:*

15: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US11A_PUBCOMB.seq:*

16: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US11A_PUBCOMB.seq:*

16: /EMC_Celerra_SIDS3/ptodata/2/pubpna/US11A_PUBCOMB.seq:*
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Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
714 3
                                                                        ID
   US-09-726-258-24
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2681.378 Million cell updates/sec
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   Sequence 24, Appl
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## ALIGNMENTS

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RESULT 1

US-09-726-24

; Sequence 24, Application US/09726258

; Sequence 24, Application US/09726258

publication No. US20030021790A1

; GENERAL INFORMATION:
   APPLICANT: Genentech, Inc., Hsei, Vanessa
   APPLICANT: Heong, Steven R.
   APPLICANT: Presta, Leonard G.
   APPLICANT: Shahrokh, Zahra
   APPL
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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; TYPE: Nucleic Acid
; STRANDEDNESS: Double
; TOPOLOGY: Linear
US-09-726-258-24
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APPLICATION NUMBER: 60/094003
APPLICATION NUMBER: 60/094003
FILING DATE: 24-JUL-1998
ATTORNSY/AGENT INFORMATION:
NAME: LOVE, RICHARD B.
REGISTRATION NUMBER: 91.659
REFERENCE/DOCKET NUMBER: P108:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECAX: 650/252-9881
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: WinPatin (Ge
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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                                                                                                                                                                                                                        GlyThrLysLeuGluIleLysArgThrValAlaAlaProSerValPheIlePheProPro
                                                                                                                                                                                                                                                                                         GluAspLeuAlaAspPhePheCysGlnGlnTyrSerAsnTyrProTrpThrPheGlyGly 100
                                                                                                                                                                                                                                                                                                                              CGCTTCACAGGCAGTGGATCTGGGGACAGATTTCACTCTCACCATCAGCCATGTGCAGTCT
                                                                                                                                                                                                                                                                                                                                                ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer
                                                 GluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThr
                                                                                                                                                  TCTGATGAGCAGTTGAAATCTGGGAACTGCTTCTGTTGTGTGCCTGCTGAATAACTTCTAT
                                                                                                                                                                        SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr
                                                                                                                                                                                                                                                                      GAAGACTTGGCAGACTATTTCTGTCAGCAATATAACATCTATCCTCTCACGTTCGGTCCT
LeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCysGluValThrHisGlnGly
                                GAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGACG
                                                                                         CCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAG
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1017.00
96.3%
90.7%
91.1%
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Matches:
Conservative:
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APPLICANT: Koumenis, Inc., Hsei, Vanessa
APPLICANT: Koumenis, Iphigenia
APPLICANT: Leong, Steven R.
APPLICANT: Leong, Steven R.
APPLICANT: Shahrokh, Zahra
APPLICANT: Shahrokh, Zahra
APPLICANT: Shahrokh, Zahra
CURRENT: Sapata, Gerardo A.
FILE REFERENCE: P1085R6
FILE REFERENCE: P1085R6
CURRENT APPLICATION NUMBER: US/11/259,232
CURRENT FILING DATE: 2005-10-25
PRIOR APPLICATION NUMBER: US/09/489,394
PRIOR APPLICATION NUMBER: US/09/489,394
PRIOR APPLICATION NUMBER: US 60/116,787
PRIOR FILING DATE: 1090-01-21
NUMBER: OF SEQ ID NOS: 72
SEQ ID NO 24
LENGUTH: 714
TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local Similarity:
Query Match:
DB:
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; NAME/KEY: Artificial Sequence
; LOCATION: 1-714
; OTHER INFORMATION: recombinant
US-11-259-232-24
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US-11-259-232-24
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                                                                                                                                       81 GluaspLeualaaspPhePheCysGlnGlnTyrSerAsnTyrProTrpThrPheGlyGly 100
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SerAspGluGlnLeuLysSerGlyThrAlaSerValValCysLeuLeuAsnAsnPheTyr
                                                                                GlyThrLysLeuGluIleLysArgThrValAlaAlaProSerValPheIlePheProPro
                                                                                                                        GAAGACTTGGCAGACTATTTCTGTCAGCAATATAACATCTATCCTCTCACGTTCGGTCCT
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No. US20060083683A1
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Matches:
Conservative:
Mismatches:
Indels:
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APPLICANT: KORDPEN, HARTMUT
APPLICANT: BOLSON, ANDREW
APPLICANT: SMITH, VICTORIA
TITLE OF INVENTION: Compositions and Methods for the Treatment of T
TITLE OF INVENTION: Compositions and Methods for the Treatment of T
TITLE OF INVENTION: Hematopoietic Origin
FILE REFERENCE: P5109R1-US
CURRENT FILING DATE: 2004-12-21
PRIOR APPLICATION NUMBER: US/11/020,508
CURRENT FILING DATE: 2004-11-16
PRIOR APPLICATION NUMBER: US 10/989,826
PRIOR FILING DATE: 2004-11-16
PRIOR APPLICATION NUMBER: US 60/520,842
PRIOR FILING DATE: 2003-11-17
PRIOR APPLICATION NUMBER: US 10/712,892
PRIOR FILING DATE: 2003-11-13
PRIOR APPLICATION NUMBER: US 10/643,795
PRIOR FILING DATE: 2003-08-19
PRIOR APPLICATION NUMBER: US 10/643,795
PRIOR APPLICATION NUMBER: US 10/643,795
PRIOR APPLICATION NUMBER: US 10/643,795
PRIOR FILING DATE: 2003-08-19
PRIOR FILING DATE: 2003-08-19
PRIOR FILING DATE: 2003-08-19
PRIOR FILING DATE: 2003-08-19
PRIOR FILING DATE: 2003-09-11
PRIOR FILING DATE: 2002-09-11
PRIOR APPLICATION NUMBER: US 10/241,220
PRIOR FILING DATE: 2002-09-11
PRIOR FILING DATE
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                                                                          Percent Similarity:
Best Local Similarity:
Query Match:
                                                                                                                                                                                 Alignment Scores: Pred. No.:
US-10-635-908-15 (1-214) x US-11-020-508-14 (1-699)
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APPLICANT:
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                                                                                                                                                                                                                                                                                     LENGTH: 699
TYPE: DNA
ORGANISM: Homo
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EBENS, ALLEN JR.
DE SAUVAGE. FREDI
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HONGO, JO-ANNE
                                                    2.16e-113
977.50
94.4%
88.8%
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Matches:
Conservative:
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Indels:
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APPLICANT: Webb, Iain J.

APPLICANT: Webb, Iain J.

APPLICANT: Horvath, Christopher J.

APPLICANT: HORVATH, Christopher J.

TITLE OF INVENTION: MCDIFIED ANTIBODIES TO PROSTATE-SPECIFIC

TITLE OF INVENTION: MEMBRANE ANTIGEN AND USES THEREOF

FILE REFERENCE: 10448-163005

CURRENT APPLICATION NUMBER: US/11/218,813

CURRENT FILING DATE: 2005-09-02

PRIOR APPLICATION NUMBER: PCT/US2004/006543

PRIOR FILING DATE: 2004-03-03

NUMBER OF SEQ ID NOS: 144

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 129
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; FEATURE:
; NAME/KEY: CDS
; LOCATION: (151)...(849)
US-11-218-813-129
                                                                                                                                                                                                                                                                                                      Sequence 129, Application US/11218813
Publication No. US20060062793A1
GENERAL INFORMATION:
                                                      LENGTH: 862
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           475
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 235 CGCTTCACTGGCAGTGGATCTGGGACGGATTTCACTTTCACCATCAGCAGTGTGCAGGCT
                                                                                                                                                                                                                                                                                                                                                                                               595
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  295 GAAGACCTGGCATTTTATTTCTGTCAGCAACATTATAGTACTCCATTCACGTTCGGCTCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer
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APPLICANT: Chisholm, Vanessa
APPLICANT: Crowley, Craig W.
APPLICANT: Krummen, Lynne A.
APPLICANT: Meng, Yu-Ju G.
APPLICANT: Meng, Yu-Ju G.
TITLE OF INVENTION: EXPRESSION VECTORS AND METILE REPERENCE: P1746R1P1 US
CURRENT APPLICATION NUMBER: US/10/714,000
CURRENT FILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: US 10/019,586
PRIOR FILING DATE: 2001-12-20
PRIOR APPLICATION NUMBER: PCT/US00/18841
PRIOR APPLICATION NUMBER: US 60/143,360
PRIOR FILING DATE: 1099-07-12
NUMBER OF SEQ ID NOS: 17
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US-10-714-000-16
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; SEQ ID NO 16
; LENGTH: 12514
; TYPE: DNA
; ORGANISM: Artificial sequence
; FEATURE:
; FEATURE:
; OTHER INFORMATION: Plasmid pSV.IPD.2C 4 circular ds-DNA
US-10-714-000-16
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Sequence 4, Application US/10715270
Publication No. US20050019925A1
GENERAL INFORMATION:
APPLICANT: Krummen, Lynne A.
APPLICANT: Shen, Amy Y.
APPLICAN
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LENGTH: 12514
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PRIOR APPLICATION NUMBER: US 60/426,095
PRIOR FILING DATE: 2002-11-14
NUMBER OF SEQ ID NOS: 6
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NAME/KEY: Misc-feature
LOCATION: 1079-1643
OTHER INFORMATION: DHFR
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OTHER INFORMATION: start
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OTHER INFORMATION: plasmid pSV.IPD.2C4 circular ds-DNA
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OTHER INFORMATION: splice
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 CCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAG 4690
                  ProArgGluAlaLysValGlnTrpLysValAspAsnAlaLeuGlnSerGlyAsnSerGln 160
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; SEQ ID NO 15
; LENGTH: 702
; TYPE: DNA
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (11..(699)
US-10-377-121-15
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Publication No. US20040001825A1

APPLICANT: GOVINDAM, SERENGULAM

APPLICANT: GOVINDAM, SERENGULAM

APPLICANT: GU, ZHENGXING

APPLICANT: HANSEN, HANS

APPLICANT: GOLDENBERG, DAVID

ITITLE OF INVENTION: RS7 ANTIBODIES

FILE REFERENCE: 018733/1163

CURRENT APPLICATION UNMER: US/10/377,121

CURRENT FILING DATE: 2003-03-03

PRIOR APPLICATION NUMBER: 60/360,299

PRIOR TILING DATE: 2002-03-01

PRIOR FILING DATE: 2002-03-01

NUMBER OF SEQ ID NOS: 36

SOPTWARE: Patentin Ver. 2.1
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Best Local Similarity:
Query Match:
DB:
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             GGRARGCCCTRAGCTCCTGATCTACTCGGCATCCTACCGGTACACTGGAGTCCCTGAT
GGGACCAAGGTGGAGATCAAACGTACTGTGGCTGCACCATCTGTCTTCATCTTCCCGCCA
                                                GluAspLeuAlaAspPhePheCysGlnGlnTyrSerAsnTyrProTrpThrPheGlyGly 100
                                                                                                 ArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIleSerAsnMetGlnSer
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Conservative:
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Sequence 19, Application US/10377121
Publication No. US20040001825A1
GENERAL INFORMATION:
APPLICANT: GOVINDAM, SERENGULAM
APPLICANT: QU. ZHENGXING
APPLICANT: HANSEN, HANS
APPLICANT: HANSEN, HANS
APPLICANT: HANSEN, HANS
CURRENT FILICATION UNMBER: US/10/377,121
CURRENT APPLICATION UNMBER: US/10/377,121
CURRENT FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: 60/360,299
PRIOR TILING DATE: 2002-03-01
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PATENTIN Ver. 2.1
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TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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                              IleThrCysLysAlaSerGlnAsnValValSerAlaValAlaTrpTyrGlnGlnLysPro
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Sequence 55, Application US/10281479A
Publication No. US20030133932A1
GENERAL INFORMATION:
APPLICANT: The UAB Research Foundation
APPLICANT: Zhou, Tong
APPLICANT: Zhou, Tong
APPLICANT: Inhikawa, Kimihisa
APPLICANT: Kimberly, Robert P.
APPLICANT: Koppman, William J.
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SOFTWARE: Patentin version 3.0
SEQ ID NO 55
LENGTH: 711
TYPE: DNA
ORGANISM: artificial sequence
FEATURE:
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Percent Similarity:
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                                                                                                          Alignment Scores:
                                                                                                                                              OTHER INFORMATION: Description US-10-281-479A-55
                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/281,479A
CURRENT FILING DATE: 2003-01-28
PRIOR APPLICATION NUMBER: 60/391,478
PRIOR FILING DATE: 2002-06-24
PRIOR APPLICATION NUMBER: 60/346,402
PRIOR FILING DATE: 2001-11-01
PRIOR APPLICATION NUMBER: PCT/US01/14151
PRIOR APPLICATION NUMBER: 60/201,344
PRIOR FILING DATE: 2001-05-02
PRIOR APPLICATION NUMBER: 60/201,344
PRIOR APPLICATION NUMBER: 60/201,344
PRIOR FILING DATE: 2000-05-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Oshumi, Jun
APPLICANT: LoBuglio, Albert S.
APPLICANT: Buchsbaum, Donald J.
TITLE OF INVENTION: COMBINATIONS OF
TITLE OF INVENTION: FACTOR-RELATED
TITLE OF INVENTION: AGENTS
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Oshumi, Jun
                1.84e-110
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                                                                                                                                                                 Artificial Sequence:/No. US20030133932Ale - Synth
                    Length:
Matches:
Conservative:
Mismatches:
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Sequence 55, Application US/10275180A

Sequence 55, Application US/10275180A

Publication No. US20030190887A1

Publication No. US20030190887A1

Publicant: The UAB Research Foundation

APPLICANT: The UAB Research Foundation

APPLICANT: Ichikawa, Kimihisa

APPLICANT: Kimberly, Robert P.

APPLICANT: Kopman, William J.

TITLE OF INVENTION: AN ANTIBODY SELECTIVE FOR A TUMOR NECROSIS FACTOR-RELATED APOPTOS

TITLE OF INVENTION: INDUCING LIGAND RECEPTOR AND USES THEREOF

FILE REFERENCE: 21085.0029U5

CURRENT APPLICATION NUMBER: US/10/275,180A

CURRENT FILING DATE: 2002-10-31

NUMBER OF SEQ ID NOS: 102

SOFTWARE: PatentIn version 3.0

SEQ ID NO 55

LENGTH: 711
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                                              OTHER INFORMATION:
OTHER INFORMATION:
US-10-275-180A-55
    Alignment Scores: Pred. No.:
                                                                                              ORGANISM: Artificial Sequence FEATURE:
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                                                              Description of Artificial Sequence:/No. US20030190687Ale Synthetic Construct
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APPLICANT: Kimberly, Robert P.
APPLICANT: Koopman, William J.
APPLICANT: LOBUGIIO, Albert S.
APPLICANT: LOBUGIIO, ALBERT SELECTIVE FOR A TUMOR NECROSIS FACTOR-RELATED TITLE OF INVENTION: APOPTOSIS-INDUCING LIGAND RECEPTOR AND USES THEREOF FILE REFERENCE: 21085.0029U7
CURRENT FILING DATE: 2003-01-22
PRIOR APPLICATION NUMBER: US 60/346,402
PRIOR APPLICATION NUMBER: US 60/301/344
PRIOR FILING DATE: 2001-05-02
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APPLICANT: Zhou, Ton
APPLICANT: Kimberly
APPLICANT: Koopman,
APPLICANT: LoBuglio
APPLICANT: Buchsbau
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SOFTWARE: PatentIn version
SEQ ID NO 55
LENGTH: 711
Sequence 127, Application US/09949559
Patent No. US20020151682A1
GENERAL INFORMATION:
APPLICANT: Athwal, Diljeet Singh
APPLICANT: Brown, Derek Thomas
APPLICANT: Weir, Andrew Neil Charles
APPLICANT: Popplewell, Andrew George
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APPLICANT: King, David John
TITLE OF INVENTION: Biological Products
ELLE REFERENCE: CAIP-0095
CURRENT FILING DATE: 2001-12-20
PRIOR APPLICATION NUMBER: 0013810.7GB
PRIOR APPLICATION NUMBER: 0013810.7GB
PRIOR FILING DATE: 2000-06-06
PRIOR FILING DATE: 2001-06-06
PRIOR FILING DATE: 2001-06-06
PRIOR FILING DATE: 2001-06-06
NUMBER OF SEQ ID NOS: 130
SOFTWARE: PALENTIN VERSION 3.1
SEQ ID NO 127
LENGTH: 642
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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RESULT 13 US-09-875-221A-127

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APPLICANT: King, David John
TITLE OF INVENTION: Biological Products
FILE REFERENCE: Carp-0089
CURRENT APPLICATION NUMBER: US/09/875,221A
CURRENT FILING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: GB0013810.7
PRIOR FILING DATE: 2000-06-06
NUMBER OF SEO ID NOS: 130
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APPLICANT: Brown, Derek Thomas
APPLICANT: Weir, Andrew Neil Charles
APPLICANT: Popplewell, Andrew George
APPLICANT: Chapman, Andrew Paul
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ORGANISM: Artificial Sequence
FEATURE:
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GAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGACG
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Sequence 3, Application US/10310454

Publication No. US20030157061A1

APPLICANT: Pharmacia Corporation

FITTLE OF INVENTION: Combinations of a Cyclooxygenase-2 Selective

FITTLE OF INVENTION: Antagonist and Therapeutic Uses Therefor

FILE REFERENCE: 18438/09008

CURRENT APPLICATION NUMBER: US/10/310,454

CURRENT FILING DATE: 2002-12-05

NUMBER OF SEQ ID NOS: 6

SOFTWARE: PatentIn version 3.1

SEQ ID NO 3

LENGTH: 642

TYPE: DNA

ORGANISM: Artificial sequence

FEATURE:

PUBLICATION OF THE SEQUENCE OF The Condition o
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; LOCATION: (1)..(642)
; OTHER INFORMATION:
US-10-310-454-3
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FILE REFERENCE: L101564WI
CURRENT APPLICATION NUMBER: US/10/401,344
CURRENT FILING DATE: 2003-03-27
NUMBER OF SEQ ID NOS: 22
SOFTWARE: PatentIn version 3.1
SEQ ID NO 3
LENGTH: 708
TYPE: DNA
ORGANISM: Homo sapiens
US-10-401-344-3
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Best Local Similarity:
Query Match:
DB:
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US-10-401-344-3
JSequence 3, Application US/10401344
Sequence 3, Application No. US20030194404A1
Publication No. US20030194404A1
GENERAL INFORMATION:
APPLICANT: Schering Corporation and Abgenix, Inc.
APPLICANT: Greenfeder, Scott
APPLICANT: Corvalan, Jose
TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO INTERLEUKIN-5 AND METHODS AND COMP
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 TCTGATGAGCAGTTGAAATCTGGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTAT
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           LeuSerSerProValThrLysSerPheAsnArgGlyGluCys 214
                                                                                   GluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThr 180
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Search completed: June Job time : 1481.01 secs 'n 2006, 07:32:32

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Command line parameters:

-MODEL-frame+ p2n.model -DEV=xlh
-Q=/abse/ABSSWEB spool/US10635908/runat 02062006 104226 10433/app query.fasta_1
-Q=/abse/ABSSWEB spool/US10635908/runat 02062006 104226 10433/app query.fasta_1
-DB=Published Applications NA New -QFMT=fastap -SUFFIX=rnpbn -MINMATCH=0.1
-DB=Published Applications NA New -QFMT=fastap -SUFFIX=rnpbn -MINMATCH=0.1
-LOOPCL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR SCORE=pct -THR MAX=100
-THR_MIN=0 -ALIGN=15 -MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
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Maximum DB seq length: 2000000000
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-NO MMAP -NEG SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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1. /EMC Celerra SIDS3/ptodata/2/pubpna/US09 NEW PUB.seq:*

2. /EMC_Celerra SIDS3/ptodata/2/pubpna/US06 NEW PUB.seq:*

3. /EMC_Celerra SIDS3/ptodata/2/pubpna/US07 NEW PUB.seq:*

4. /EMC_Celerra_SIDS3/ptodata/2/pubpna/US08 NEW PUB.seq:*

5. /EMC_Celerra_SIDS3/ptodata/2/pubpna/PCT NEW PUB.seq:*

6. /EMC_Celerra_SIDS3/ptodata/2/pubpna/US10 NEW PUB.seq:*

7. /EMC_Celerra_SIDS3/ptodata/2/pubpna/US10 NEW PUB.seq:*

8. /EMC_Celerra_SIDS3/ptodata/2/pubpna/US10 NEW PUB.seq:*
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Copyright (c) 1993 - 2006 Biocceleration Ltd.

    2006, 06:56:49; Search time 24.4204 Seconds
(without alignments)
    1548.108 Million cell updates/sec

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US-11-219-563-129

US-10-981-300-17

US-11-211-917-23

US-10-546-594-131

US-10-981-300-13

US-11-155-444-3

US-11-211-917-87
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Sequence 10, Appl
Sequence 129, Appl
Sequence 17, Appl
Sequence 23, Appl
Sequence 131, App
Sequence 131, Appl
Sequence 13, Appl
Sequence 47, Appl
Sequence 47, Appl
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Alignment Score pred. No.: Score: Score: Percent Similar Best Local Simi Query Match: DB:	RESULT 1  WS-11-106-762-10  Sequence 10, Ap  Publication No.  GENERAL INFORM  FILLE REFERENCE  FILLE REFERENCE  CURRENT PAPLICAN  CURRENT FILING  CURRENT FILING  PRIOR APPLICAN  PRIOR FILING I  NUMBER OF SEQ  SEQ ID NO 10  LENGTH: 1701  TYPE: DNA  ORGANISM: Mus  FEATURE:  NAME/KEY: Une  LOCATION: 166  CUTHER INFORM  US-11-106-762-10		10 878 11 12 13 869 14 869 15 867 16 867 17 865 19 865 19 865 20 865 21 865 21 865 22 863 23 843 24 869 25 689 27 649 30 624 31 51 31 32 51 33 51 34 35 47 482 482 482 482 466
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Length: Matches: Conservative: Mismatches: Indels: Gaps:	06762 ET AL. ANTIBODIES /11/106,762 15 0/563,193	ALIGNMENTS	-155-444-19 -211-917-71 -211-917-31 -231-97-1838 -211-97-1838 -211-97-1838 -211-97-197-101 -293-697-1863-1 -297-317-18 -297-31
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			19, Appl 31, Appl 31, Appl 31, Appl 31, Appl 31, Appl 15, Appl 17, Appl 55, Appl 63, Appl 63, Appl 63, Appl 64, Appl 65, Appl 66, Appl 67, Appl 68, Appl 69,

US-10-635-908-15 (1-214) x US-11-106-762-10 (1-1701)

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TITLE OF INVENTION: MCDIFIED ANTIBODIES TO PROSTATE-SPECIFIC TITLE OF INVENTION: MCMMBRANE ANTIGEN AND USES THEREOF FILE REFERENCE: 13651.001 (BZI-001)
CURRENT APPLICATION NUMBER: US/11/219,563
CURRENT FILING DATE: 2005-09-02
PRIOR APPLICATION NUMBER: PCT/US04/06586
PRIOR FILING DATE: 2004-03-03
PRIOR FILING DATE: 2004-03-03
PRIOR FILING DATE: 2004-03-03
PRIOR FILING DATE: 2003-03-03
PRIOR FILING DATE: 2003-03-03
PRIOR FILING DATE: 2003-05-30
VMMBER OF SEQ ID NOS: 144
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 129
LENGTH: 862
TYPE: DNA
   FEATURE:
COTHER INFORMATION: Light correction: Light correction: Light correction: Light correction: (151) ... (849)
US-11-219-563-129
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APPLICANT: Bander, Neil
                                                                                          ORGANISM: Artificial Sequence
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US-10-981-300-17

(Sequence 17, Application US/10981300

) Publication No. US20060093599A1

; GENERAL INFORMATION:

; APPLICANT: GIORGIO SENALIDI

; APPLICANT: GADI GAZIT-BORNSTEIN

; TITLE OF INVENTION: ANTI-PROPERDIN ANTIBODIES, ANI

; TITLE OF INVENTION: FOR MAKING AND USING THE SAMI

; FILE REFERENCE: ABGX-005

; CURRENT APPLICATION NUMBER: US/10/981,300

; CURRENT FILING DATE: 2004-11-03

; NUMBER OF SEQ ID NOS: 71

; SOFTMARE: FASTSEQ for Windows Version 4.0

; ENGTH: 642

; TYPE: DNA

; ORGANISM: homo sapien
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567 120 507 100 447 80 60 327 40 267

849 214

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Sequence 23, Application US/11211917

Publication No. US20060093600A1

GENERAL INFORMATION:
APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONALD P.
APPLICANT: GLADUE, RONALD P.
APPLICANT: JIA, XIAO-CHI
APPLICANT: JIA, XIAO-CHI
APPLICANT: JIA, XIAO-CHI
FILE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-FP/3 US
CURRENT APPLICATION NUMBER: US/11/211,917
CURRENT ETLING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR APPLICATION NUMBER: 60/348,980
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       RESULT 5
US-10-546-594-131
Sequence 131, Application US/10546594
Publication No. US20060088538A1
GENERAL INFORMATION:
APPLICANT: HOSOKAWA, Saiko
APPLICANT: HOSOKAWA, Soiko
APPLICANT: HIRAKAWA, Yoko
APPLICANT: HIRAKAWA, Yoko
APPLICANT: LTAMI, Seima
APPLICANT: UMEKI, Hiroe
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NUMBER OF SEQ ID NOS: 147
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 23
LENGTH: 705
TYPE: DNA
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US-10-635-908-15 (1-214) x US-11-211-917-23 (1-705)
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         LeuSerSerProValThrLysSerPheAsnArgGlyGluCys
                                               LeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCysGluValThrHisGlnGly
                                                                                         GluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThr
                                                                                                                     GluAspLeuAlaAspPhePheCysGlnGlnTyrSerAsnTyrProTrpThrPheGlyGly 100
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TITLE OF INVENTION: MONOCLONAL ANTIBODY AND GENE ENCODING THE SAME, HYBRIDOMA,
TITLE OF INVENTION: MONOCLONAL ANTIBODY AND DIAGNOSTIC REAGENT
FILE OF INVENTION: COMPOSITION, AND DIAGNOSTIC REAGENT
FILE REFERENCE: 238067
FILE REFERENCE: 238067
CURRENT FILING DATG: 2005-08-19
PRIOR APPLICATION NUMBER: US/10/546,594
CURRENT FILING DATG: 2003-02-28
PRIOR APPLICATION NUMBER: JP 2003/54670
PRIOR FILING DATG: 2003-07-09
PRIOR FILING DATG: 2003-07-09
NUMBER OF SEQ ID NOS: 132
SOFTWARE: Patentin version 3.1
SEQ ID NO 131
LENGTH: 705
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APPLICANT:
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LOCATION: (1)..(705)
OTHER INFORMATION:
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TYPE: DNA
ORGANISM: Homo sapiens
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                    LeuSerLy8AlaAspTyrGluLy8HisLy8ValTyrAlaCy8GluValThrHisGlnGly
                                                                 GAGAGTGTCACAGAGCAGGACAGCAAGGACACCTACAGCCTCAGCAGCACCCTGACG
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   CTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAGGGC
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KUMAI, Koichiro
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; Sequence 13, Application US/10981300
; Publication No. US20060093599A1
; GENERAL INFORMATION:
; APPLICANT: GIORGIO SENALDI
; APPLICANT: GADI GAZIT-BORNSTEIN
; TITLE OF INVENTION: ANTI-PROPERDIN ANTIBODIES, AND MET
; TITLE OF INVENTION: FOR MAKING AND USING THE SAME
; TITLE REFERENCE: ABGX-005
; CURRENT APPLICATION NUMBER: US/10/981,300
; CURRENT FILING DATE: 2004-11-03
; NUMBER OF SEQ ID NOS: 71
; SOFTWARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 13
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; TYPE: DNA
; ORGANISM: homo 6
US-10-981-300-13
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LeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCysGluValThrHisGlnGly
                                                                     GluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeuSerSerThrLeuThr
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                                                  GAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTGACG
                                                                                                                 CCCAGAGAGGCCAAAGTACAGTGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCCAG
                                                                                                                                   ProArgGluAlaLysValGlnTrpLysValAspAsnAlaLeuGlnSerGlyAsnSerGln
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PRIOR FILING DATE: 2003-12-22
PRIOR APPLICATION NUMBER: 60/435,154
PRIOR APPLICATION NUMBER: 60/435,154
PRIOR FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: 60/435,185
PRIOR APPLICATION NUMBER: 60/435,185
PRIOR FILING DATE: 2002-12-20
NUMBER OF SEQ ID NOS: 23
SOPTWARE: Patentin Ver. 3.3
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: GARBER, ELLEN
APPLICANT: BAILLY, VERONIQUE
APPLICANT: BAILLY, VERONIQUE
APPLICANT: BROWNING, JEFFREY L.
TITLE OF INVENTION: MULTIVALENT LYMPHOTOXIN BETA RECEPTOR AGONISTS AND
TITLE OF INVENTION: THERAPEUTIC USES THEREOF
FILE REFERENCE: BGNA168CN
CURRENT APPLICATION NUMBER: US/11/155,444
CURRENT APPLICATION NUMBER: US/11/155,444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence: Light chain OTHER INFORMATION: huCBEll/huBHA10 bispecific-1 antibody construct
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               GlyThrLysLeuGluIleLysArgThrValAlaAlaProSerValPheIlePheProPro 120
                                                                                                                         GluAspLeuAlaAspPhePheCysGlnGlnTyrSerAsnTyrProTrpThrPheGlyGly 100
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; Sequence 47, Application US/11211917
; Publication No. US20060093600A1
; GENERAL INFORMATION:
; APPLICANT: BEDIAN, VAHE
; APPLICANT: GLADUE, RONALD P.
; APPLICANT: CORVALAN, JOSE
; APPLICANT: JIA, XIAO-CHI
; APPLICANT: JIA, YIAO-CHI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION UNMBER: US/11/211,917
CURRENT FILING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR FILING DATE: 2002-11-08
PRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 47
LENGTH: 705
TYPE: DNA
ORGANISM: Homo sapiens
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; NAME/KEY: CDS ; LOCATION: (1) US-11-155-444-3

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Pred. No.: Alignment Scores:

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FILE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/11/211,917
CURRENT FILING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR FILING DATE: 2002-11-08
PRIOR FILING DATE: 2002-11-09
PRIOR APPLICATION NUMBER: 60/348,980
PRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 87
LENGTH: 705
TYPE: DNA
ORGANISM: Homo sapiens
US-11-211-917-87
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; Sequence 87, Application US/11211917
; Publication No. US20060093600A1
; GENERAL INFORMATION:
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APPLICANT: GLADUE, RONALD
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
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US-11-155-444-19
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                                                                    Score:
                                                                                           Pred. No.:
                                                                                                              Alignment
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                                                                                                                                                                                                                                                    OTHER INFORMATION:
                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
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79.4%
80.6%
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APPLICANT: GARBER, ELLEN
APPLICANT: GARBER, ELLEN
APPLICANT: BAILLY, VERONIQUE
APPLICANT: BAOWNING, JEFFREY L.
TITLE OF INVENTION: THERAPEUTIC USES THEREOF
FILE OF INVENTION: THERAPEUTIC USES THEREOF
FILE REFERENCE: BGNA168CN
CURRENT APPLICATION NUMBER: US/11/155,444
CURRENT FILING DATE: 2005-06-17
PRIOR APPLICATION NUMBER: PCT/US03/041393
PRIOR FILING DATE: 2003-12-22
PRIOR APPLICATION NUMBER: 60/435,154
PRIOR FILING DATE: 2003-12-20
PRIOR APPLICATION NUMBER: 60/435,185
PRIOR APPLICATION NUMBER: 60/435,185
PRIOR APPLICATION NUMBER: 60/435,185
PRIOR FILING DATE: 2002-12-20
PRIOR APPLICATION ONDER: 60/435,185
PRIOR FILING DATE: 2002-12-20
SAUMBER OF SEQ ID NOS: 23
SOFTWARE: PATENTIN Ver. 3.3
SEQ ID NO 19
LENGTH: 645
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                                                                                                                                                                                                                                                                                     Description of Artificial Sequence: CBE11 chimeric light chain antibody construct
                     Length:
Matches:
Conservative:
Mismatches:
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Sequence 71, Application US/11211917
Sequence 71, Application US/20060093600A1
GENERAL INFORMATION:
APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONALD P.
APPLICANT: GLADUE, RONALD P.
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: TIA, XIAO-CHI
OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT FILING DATE: 2005-08-25
CURRENT FILING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR FILING DATE: 2002-11-08
PRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: PATENTIN VET: 2.1
SEQ ID NO 71
LENGTH: 705
TYPE: DNA
ORGANISM: Homo sapiens
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Sequence 39, Application US/11211917

Publication No. US20060093600A1

GENERAL INFORMATION:
APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONALD P.
APPLICANT: CORVALAN, JOSE
APPLICANT: CORVALAN, JOSE
APPLICANT: FENG, XIAO-CHI
APPLICANT: FENG, XIAO-CHI
APPLICANT: FENG, XIAO-CHI
CURRENT APPLICATION ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/11/211,917
CURRENT FILING DATE: 2005-10/292,088
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR APPLICATION NUMBER: 60/348,980
PRIOR APPLICATION NUMBER: 60/348,980
PRIOR FILING DATE: 2001-11-09

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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
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Pred. No.:
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RESULT 12
US-11-211-917-39
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                                GlyLeuSerSerProValThrLysSerPheAsnArgGlyGluCys 214
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                                                                                                                          CAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTG
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                                                                           ACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCCATCAG
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; NUMBER OF SEQ ID NOS
; SOFTWARE: Patentin V
; SEQ ID NO 39
; LENGTH: 720
; TYPE: DNA
; ORGANISM: Homo Bapi
US-11-211-917-39
Sequence 31, Application US/11211917
Publication No. US20060093600A1
GENERAL INFORMATION:
APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONALD P.
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
TITLE OF INVENTION: ANTIBODIES TO CD4
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Best Local Similarity:
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; TYPE: DNA
; ORGANISM: Homo &
US-11-211-917-31
RESULT 14
US-11-293-697-1838
; Sequence 1838, Application US/11293697
; Publication No. US20060105376A1
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Best Local Similarity:
Query Match:
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FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/11/211,917
CURRENT FILING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR FILING DATE: 2002-11-08
PRIOR APPLICATION NUMBER: 60/348,980
PRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: PATENTIN Ver. 2.1
SOFTWARE: PATENTIN Ver. 2.1
SOFTWARE: PATENTIN Ver. 2.1
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ValThrHisGlnGlyLeuSerSerProValThrLysSerPheAsnArgGlyGluCys 214
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                                                                                                    AGCAGCACCCTGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA
                                                                                                                            SerSerThrLeuThrLeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCysGlu
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GENERAL INFORMATION:
APPLICANT: HELIX RESEARCH INSTITUTE
TITLE OF INVENTION: NOVEL full length cDNA
FILE REFERENCE: H1-A0106
CURRENT APPLICATION NUMBER: US/11/293,697
CURRENT FILING DATE: 2005-12-05
PRIOR APPLICATION NUMBER: US/10/108,260
PRIOR APPLICATION UNMER: US/10/108,260
PRIOR FILING DATE: 2002-03-28
NUMBER OF SEQ ID NOS: 5458
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 1838
LENGTH: 929
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Best Local Similarity:
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                                                           LeuSerSerProValThrLysSerPheAsnArgGlyGluCys
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 Application US/11211917
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GENERAL INFORMATION:
APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONALD P.
APPLICANT: GLADUE, RONALD P.
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
TITLE OF INVENTION: ANTIBODIES TO CD40
TITLE REFERENCE: ABX-PF/3 US
CURRENT FILING DATE: 2005-08-25
CURRENT FILING DATE: 2005-08-25
PRIOR APPLICATION NUMBER: US/11/211,917
CURRENT FILING DATE: 2002-11-08
PRIOR APPLICATION NUMBER: US/10/292,088
PRIOR APPLICATION NUMBER: 00/348,980
PRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
NUMBER OF SEQ ID NOS: 147
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; ORGANISM: Homo sapiens US-11-211-917-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-635-908-15 (1-214) x US-11-211-917-15 (1-720)
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SEQ ID NO 15
LENGTH: 720
TYPE: DNA
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                                                                                                                                                                                                                                 116
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                                                                                                                                                                                                                                                                                                                                                                                                      241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     36 TyrGlnGlnLysProGlyGlnSerProLysLeuLeuIleTyrSerAlaSerAsnArgTyr 55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          121
                                                                                                                                   481
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601
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 GATATTGTGATGACTCAGTCTCCACTCTCCCTGCCCGTCACCCCTGGAGAGCCGGGCCTCC
                                                                                                                                                    LeuAsnAsnPheTyrProArgGluAlaLysValGlnTrpLysValAspAsnAlaLeuGln
                                                                                                                                                                                                  TCCGGGGTCCCTGACAGGTTCAGTGGCAGTGGATCAGGCACAGATTTTACACTGAAAATC
                                                                                                                                                                                                                                                                                                                                                                                                                       ThrGlyValProAspArgPheThrGlySerGlySerGlyThrAspPheThrLeuThrIle 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AspIleValMetThrGlnSerGlnArgPheMetSerThrThrValGlyAspArgValSer 20
                                                                                                                                                                                                                                                                  CGGACGTTCGGCCAAGGGACCAAGGTGGAAATCAAACGAACTGTGGCTGCACCATCTGTC
                                                                                                                                                                                                                                                                                    TrpThrPheGlyGlyGlyThrLysLeuGluIleLysArgThrValAlaAlaProSerVal 115
                                                                                                                                                                                                                                                                                                                                                                   SerAsnMetGlnSerGluAspLeuAlaAspPhePheCysGlnGlnTyrSerAsnTyrPro
                   SerSerThrLeuThrLeuSerLysAlaAspTyrGluLysHisLysValTyrAlaCysGlu
                                                                                   SerGlyAsnSerGlnGluSerValThrGluGlnAspSerLysAspSerThrTyrSerLeu
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 AGCAGCACCCTGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAA
                                                                TCGGGTAACTCCCAGGAGAGTGTCACAGAGCAGGACAGCAAGGACAGCACCTACAGCCTC
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867.50
86.3%
77.2%
77.7%
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Matches:
Conservative:
Mismatches:
Indels:
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